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McKay,R. Dean,N., Monia,B.P., Nero,P.: Antisense oligonucleotide compositions modulation of JNK proteins Patent: US 6133246-A 42 17-OCT-2000; Location/Qualifiers 120 /organism="unknown"	ARI16461 Sequence 42 from patent US ARI16461 ARI16461.1 GI:14096783 Unknown. Unknown. Unclassified.	1 GTCCGGGCCAGGCCAAAGTC 20 1 GTCCGGGCCAGGCCAAAGTC 20 1 GTCCCGGGCCAGGCCAAAGTC 20 1 GTCCCGGGCCAGGCCAAAGTC 20	Unknown: Unclassified. 1 (bases I to 20) McKay,R., Dean,N., Monia,F McKay,R., Monia,F McKay,R	aR116450 AR116450 AR116450 AR116450 AR116450 AR116450 AR116470 AR116470 AR116470 AR116470	HΩ	Sequence 7 from patent US ARI10470 ARI10470.1 GI:12826746 . Unknown. Unknown. Unclassified. 1 (bases 1 to 20) Monia,B.P. and Xu,X.S. Methods of modulating tumc expression of cell adhesic Patent: US 6114517-A 7 055
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PN JP 2001514905-A/31
PD 18-SEP-2001
PF 07-AUG-1998 JP 2000509875
PR 13-AUG-1997 US 08/910629
PI ROBERT MCKAY,NICHOLAS DEAN,BRETT P MCNILLIAM A GAARDE
PC C1201/68,A61K31/7088,A61K48/00,A61P35
PC C1201/68,A61K31/7088,A61K48/00,A61P35
PC C12N15/00
CC antisense sequence
FH Key 1. 20
FT source /organism='Artificial
                  PC CC FT
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Mckay,R., Dean,N., Monia,B.P.,
Antisence oligonucleotide comp
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synthetic construct
artificial sequences.
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C12N15/00
                                                                                                                                                                               PHARMACEUTICALS INC
Artificial Sequence
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13-AUG-1997 US 08/910629
ROBERT MCKAY,NICHOLAS DEAN,BRETT P MONTA,PAMELA SCOTT
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13-AUG-1997 US 08/910629
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/organism='Artificial Sequence'
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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                                                                                        Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J. Method and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6346398-A 6244 12-FEB-2002;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; l
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/mol_type="genomic DNA
/db_xref="taxon:32630"
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/db_xref="taxon:9606"
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Pavco.P., McSwiggen,J., Stinchcomb,D. and Escobedo,J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398,A 7661 12-FEB-2002;
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Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J. Method and reagent for the treatment of diseases or crelated to levels of vascular endothelial growth fact Patent: US 6346398-A 4222 12-FEB-2002;
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Pavco,P., McSwiggen,J., Stinchcomb,D. and Escobedo,J.
Method and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6346398-A 1996 12-FEB-2002;
                                                                                                                                                                                                                                    Sequence
AR186508
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Elledge, S.J. and Sanchez, Y.
Elledge, S.J. and Sanchez, Y.
Mammalian checkpoint genes and proteins
Patent: US 6218109-A 6 17-APR-2001;
Location/Qualifiers
                                                                                                                                                                                          Unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                          1 (bases 1 to 17)
Elledge, S. J. and Sanchez, Y.
Mammalian checkpoint genes and proteins
Patent: US 6307015-A 6 23 OCT-2001;
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AR190307
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 (bases 1 to 17)
Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 634639-A 7660 12-FEB-2002;
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AR192172
Method of amplifying DNA fragment, apparatus for amplifying DNA
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Pavco, P., McSwiggen, J.,
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                Inoue, T.
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OS Unidentified
PN JP 1999276176-A/100
PD 12-CCT-1999
PF 31-MAR-1998 JP 199808
PR KOICHI INOUE
PC C12N15/09, B09B3/00, C1:
Strandedness: Single;
FH Key
FT Source
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FOR COUNTY OF THE COUNTY OF 
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Method and device for a
E38726
E38726.1 GI:18621388
JP 200027087-A/100.
                    unidentified unclassified.
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microorganism existing and method f
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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OS Unidentified
PN JP 2000270867-A/100
PD 03-CCT-2000
PF 19-MAR-1999 JP 1999
PR KOICHI INOUE
PC C12M15/09, C12M1/00,
CC Strandedness: Singla
CC Topology: Linear;
FH Key
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FT T
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Method for amplifying DNA fragment, amplification apparatus of DNA fragment, method for assaying a group of microorganisms, method for analyzing a group of microorganisms, and method for assaying substance.
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                                                                                                                                                                                                                                         SANYO ELECTRIC CO LTD, SO
FORESTRY AND FISHERIES
OS Artificial Sequence
PN JP 1999341989-A/100
                                                                                                                                                                                                                                                                                      Method for amplifying DNA fragment, amplification apparatus of DNA fragment, method for assaying a group of microorganisms, method for analyzing a group of microorganisms, and method for assaying contaminating substance contaminating substance Patent: JP 199341989-A 100 14-DEC-1999; SANYO ELECTRIC CO LTD, SOCIETY FOR TECHNO-INNOVATION OF AGRICULTURE SANYO ELECTRIC CO LTD, SOCIETY FOR TECHNO-INNOVATION OF AGRICULTURE
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                                                                                                  artificial sequences.
1 (bases 1 to 12)
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JP 1999341989-A/100.
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                                                                           /organism='Artificial Sequence'
           /organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon;32630"
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/mol_type="genomic_DNA"
/db_xref="taxon:32644"
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Patent: WO 0138577-A 1410 31-MAY-2001;
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                                                                                                                                                             Human transcriptomes
Patent: WO 0138577-A 1411 31-MAY-2001;
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                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/mol_type="genomic DNA"
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Sequence 77
AR136587
AR136587.1
                          Unclassified.

1 (bases 1 to 13)
Andersson,M.K., Berglund,L.G.T., Reneland,R.H.
Andersson,M.K., Berglund,L.G.T., Reneland,R.H.
Reagents and methods for detection of heart fai
Patent: US 6528268-A 36 04-MAR-2003;
Location/Qualifiers
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Andersson,M.K., Berglund,L.G.T., Reneland,R.H. and Adam,G.I.R.
Reagents and methods for detection of heart failure
Patent: US 6528268-A 11 04-MAR-2003;
Location/Qualifiers
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Li,L. and Hood,L.
Human jagged polypeptide, encoding nucleic Patent: US 6136952-A 77 24-OCT-2000;
Location/Qualifiers
                                                                                             Unknown.
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                                                                                                                              GI:29722030
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1 (Dases 1 to 13)

S Gillies, S. D., Lo, K.M. and Lan, Y.

Hetero dimer fused proteins useful against targeting imm and general immunostimulation

L Patent: JP 2001525423-A 7 11-DEC-2001;

LEXIGEN PHARMACEUTICALS CORP
OS Artificial Sequence
PN P2001525423-A/7

PD 11-DEC-2001

PR 08-DEC-1998 UP 2000524321

PC C07K19/00, A61K47/48, A61P37/04, C07K14/54, C07K16/30, CC

PC C12P21/00/, A61K47/48, A61P37/04, C07K14/54, C07K16/30, CC

PC A61K38/00, C12N15/00, A61K37/02

CC Description of Artificial Sequence: Synthetic Oligon

FT source

/organism='Artificial Sequence'.
                                                                                                                           AR301743
Sequence 32
AR301743
AR301743.1
Compositions and methods for wound healing Patent: US 6538173-A 324 25-MAR-2003; Location/Qualifiers
                                                                                                                                                                                                                                                            3 CCGGGCCAGGCCA 15
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                                         Heber-Katz, E.
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08-DEC-1997 US 08/986597
STEPHEN D GILLIES, KIN MING LO, YAN LAN
C07K19/00, A61K47/48, A61P37/04, C07K14/54, C07K16/30, C12N15/09,
C12P21/00//
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Location/Qualifiers
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Sequence 4426 from Patent
AX627385
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Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Method for determining homeostasis of the skin
Patent: WO 02053774-A 166 11-JUL-2002;
Henkel Kommanditgesellschaft auf Aktien (DE)
                                                                                                                                                        Method for determining homeostasis of the sk
Patent: WO 02053774-A 4426 11-JUL-2002;
Henkel Kommanditgesellschaft auf Aktien (DE)
                                                                                                                                                                                               Petersohn,D., Conradt,M. and Hofmann,K.
                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                 AX627385.1 GI:28455423
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                   /mol_type="genomic DNA"
/db_xref="taxon:9606"
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OS MUS musculus (mou
PN JP 2002503460-A/2
PD 05-FEB-2002
PF 12-FEB-1999 JP 20
PR 13-FEB-1998 US
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Sequence 7587 from Patent
AX630546
AX630546.1 GI:28458584
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
1, (bases 1 to 11)
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PI ELLEN HEBI
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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13-FEB-1998 US 60/074737,26-AUG-1998
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AX482038/c
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RESULT 36
AX623635/c
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                                                                                                                                                                                                                                                                                         synthetic construct
synthetic construct
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Sequence 15 from Patent WO02059558.
AX511277
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Means and methods for treatment evaluation
Patent: EP 122523-A 15 24-JUL-2002;
Amsterdam Support Diagnostics B.V. (NL)
Location/Qualifiers
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artificial sequences.
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/db_xref="taxon:32630"
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/db_xref="taxon:32630"
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AX625382/c
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Sequence 2423 from Patent
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AX625382.1 GI:28453323
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Sequence 676 from Patent
AX623635
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Sequence 6380 from Patent
AX629339
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Patent: WO 02053774-A 2423 11-JUL-2002;
Henkel Kommanditgesellschaft auf Aktien (DE)
                                       Method for determining homeostasis of the skin Patent: WO 02053774-A 6380 11-JUL-2002; Henkel Kommanditgesellschaft auf Aktien (DE) Location/Qualifiers
                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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organism="Homo sapiens"
/mol_type="genomic DNA"
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AX631056/c
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Homo sapiens
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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                                                                                  /organism="Homo sapiens"
/mol_type="genomic DNA"
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AX152505/c
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Sequence 420 from Patent
AX152505
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Sequence 421 from Patent
AX152506
AX152506.1 GI:14534157
Homo sapiens
Homo sapiens
                                                                                                                                                                                                                                                  Patent: WO 0138577-A 420 31-MAY-2001;
The Johns Hopkins University (US)
Location/Qualifiers
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Lopez-Nieto, C.Eduardo. and Nigam, S.Kumar.
Processes, apparatus and compositions for characterizing nucleotide sequences based on K-tuple analysis
Patent: US 6110667-A 63 29-AUG-2000;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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Sequence
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Human transcriptomes
Patent: WO 0138577-A 644 31-MAY-2001;
The Johns Hopkins University (US)
Location/Qualifiers
                                                                                                                                                                                  Sequence 644 from Patent AX152729
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Patent: WO 0138577-A 422 31-WAY-2001;
The Johns Hopkins University (US)
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Patent: WO 0138577-A 421 31-MAY-2001;
The Johns Hopkins University (US)
                                                           Velculescu, V.E., Vogelstein, B.
                                                                                         Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Sequence 1228 from Patent WO0138577.
AX153313
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Patent: WO 0138577-A 1229 31-MAY-2001;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Patent: WO 0138577-A 1228 31-MAY-2001;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/mol_type="genomic DNA"
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Sequence 1232 from Patent WO0138577.
                                                                                                                            Similarity
9; Conserv
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Sequence 1231 from Patent
AX153316
AX153316.1 GI:14534967
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Patent: WO 0138577-A 1230 31-MAY-2001;
The Johns Hopkins University (US)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/db_xref="taxon:9606"
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/mol type="genomic DNA"
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Sequence 1233 from Pate
AX153318
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Velculescu, V.E., Vogelstein, B. and Kinzler, K.W.
                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae
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Patent: WO 0138577-A 1233 31-MAY-2001;
The Johns Hopkins University (US)
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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The Johns Hopkins University (US)
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/mol_type="genomic DNA"
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AX224404
                                              Albertsen, M.C., Fox, T.W., Garnaat, C.W., Huffman, G. Male tissue-preferred regulatory region and method Patent: WO 0160997-A 14 23-AUG-2001; PIONEER HI-BRED INTERNATIONAL, INC. (US)
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                                                                                                                           Patent: WO 0185941-A 194 15-NOV-2001;
Academisch Ziekenhuis bij de Universiteit
Location/Qualifiers
                                                                                                                                                                                                                          Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea.
                                                                                                                                                                           Versteeg, R. and Caron, H.N.
                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
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                                                                   /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
5 c 3 g
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/db_xref="taxon:4577"
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Patent WO0160997.
                      Score 8.4; DB Pred. No. 51; 0; Mismatches
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Pred. No. 51;
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Catarrhini; Hominidae;
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SOURCE
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BD065145/c
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Best Local S
Matches 9
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TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JOURNAL
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LPS activated human monocyte expressing gen

Patent: JP 2001069993-A 255 21-MAR-2001;

JAPAN SCIENCE AND TECHNOLOGY CORP

OS Homo sapiens (human)

PN JP 2001069993-A/255

PD 21-MAR-2001 JP 2000131079

PR 28-APR-2000 JP 2000131079

PR
                                                Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.

1 (bases 1 to 10)

Velculescu, V.E., Vogelstein, B. and Kinzler, K.W.
Characterization of the yeast transcriptome
Patent: JP 2001509017-A 81 10-JUL-2001;
THE JOHNS UNIVERSITY SCHOOL OF MEDICINE
OS Saccharomyces cerevisiae (yeast)
PN JP 2001509017-A/81
PD 10-JUL-2001
PF 22-JAN-1998 JP 1998532117
PR 23-JAN-1997 US 60/035917
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PD 10-JUL-2001

PF 22-JAN-1998 JP 1998532117

PF 23-JAN-1997 US 60/035917

PR 23-JAN-1997 US 60/035917

PI VICTOR E VELCULESCU, BERT VOGELSTEIN, KENNETH W KINZLER PC C12N15/10, C12N15/31, C07K14/395, C12Q1/68, C12Q1/02 CC Characterization of the yeast transcriptome
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JP 2001509017-A/81
Saccharomyces cerevisiae (baker's yeast)
Saccharomyces cerevisiae
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C12N15/09,C07K14/47,C07K16/18,G01N33/50,G01N33/53//A61K45/00,
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JP 2001069993-A/255.
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Location/Qualifiers
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Nagai, S.,
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1 (bases 1 to 10)

Matcushima, K., Hashimoto, S., Suzuki, T. and Nagai, S.

Human matured/activated dendritic cell expression genes

Patent: JP 2001327293-A 45 27-NOV-2001;

JAPAN SCIENCE AND TECHNOLOGY CORP

OS Homo sapiens (human)

PN JP 2001327293-A/45

PD 27-NOV-2001 JP 2000150562

PF 22-MAY-2000 JP 2000150562
                                                                                    Human activated Th1 and BD161262 BD161262.1 GI:27867020 JP 2002186482-A/84.
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            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 10)
Nagai,S., Matsushima,K. and Hashimoto,S.
                                                                                                                                     BD161262
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22-MAY-2000 JP 2000150562
KOJI MATSUSHIMA,SHINICHI HASHIMOTO,TAKUJI SUZUKI,SHIGENORI
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/db_xref="taxon:9606"
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RESULT 63
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Best Local S
Matches 9
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TITLE
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JOURNAL
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Human liver disease-expressing genes
Patent: JP 2002209591-A 32 30-UUL-2002;
JAPAN SCIENCE AND TECHNOLOGY CORP
OS Homo sapiens (human)
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OS Homo sapiens (human)
PN JP 2002186482-A/84
PD 02-JUL-2002
PF 19-DEC-2000 JP 20003B5816
PF 19-DEC-2000 JP 20003B5816
PF SHIGENORI NAGAI, KOJI MATSUSHIMA, SHINICHI HASHIMOTO PC C12N15/09, C07K14/47, C07K16/18, C12P21/08, C1NN15/00 CC Huma activated Th1 and Th2 cell expression genes FH Key
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Patent: JP 2002186482-A 84 02-JUL-2002;
JAPAN SCIENCE AND TECHNOLOGY CORP
OS Homo sapiens (human)
PN JP 2002186482-A/84
PD 02-JUL-2002
PF 19-DEC-2000 JP 2000385816
                                                                                                                                                                                                                                                                                                     PF DG SS
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JP 2002209591-A/32.
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                                                     Conservative
                                                                                                                                                                                                                 Human liver disease-expressing genes
Key Location/Qualifiers
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30-JUL-2002
19-JAN-2001 JP 200
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KOJI MATSUSHIMA, SHINICHI
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                                                                                                      /organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
2 c 2 g
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Location/Qualifiers
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/db_xref="taxon:9606"
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/mol_type="genomic DNA"
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Pred. No. 51;
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RESULT 64

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BD167212/c
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PN JP 20
PD 30-JI
PF 19-JI
PF KOJI
YAMASHITA
PC C12NI
PC C12NI
PC C12NI
CC Humaa
CC Humaa
FH Key
FT Source
                                                                                                                                                                          unclassified.

1 (bases 1 to 10)

Matsushima, K., Hashimoto, S., Kaneko, S. and Human liver disease-expressing genes Patent: JP 2002209591-A 757 30-JUL-2002;
JAPAN SCIENCE AND TECHNOLOGY CORP OS Homo sapiens (human)
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Matsushima, K., Hashimoto, S., Kaneko, S. ar

Human liver disease-expressing genes

Patent: JP 2002209591-A 451 30-JUL-2002;

JAPAN SCIENCE AND TECHNOLOGY CORP

OS Homo sapiens (human)

PN JP 2002209591-A/451
                                                                                                                                                                                                                                                                                        JP 2002209591-A/757. unidentified
                 3336888
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                                                                                                                        19-JAN-2001 JP 2001012328
KOJI MATSUSHIMA, SHINICHI HASHIMOTO, SHUICHI KANEKO, TARO
                                       Human liver disease-expressing genes 
Key Location/Qualifiers
                                                                       C12N15/00
                                                                                  C12N15/09,C07K14/47,C07K16/18,G01N33/15,G01N33/50//C12P21/02,
C12P21/08,
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Key Location/Qualifiers
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19-JAN-2001 JP 2001012328
KOJI MATSUSHIMA,SHINICHI HASHIMOTO,SHUICHI KANEKO,TARO
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/organism='Homo sapiens (human)'.
Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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Pred. No. 51;
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KEYWORDS
SOURCE
ORGANISM
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VERSION
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E39743/c
LOCUS
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AUTHORS
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E39489/c
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Best Local S
Matches
                AUTHORS
TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                     OS Homo sapiens (human)
PN JP 2000279181-A/22
PD 10-CCT-2000
PF 01-APR-1999 JP 1999095481
PR SHINICHI HASHIMOTO, KOJI MATSUSHIMA, TAKUJI SU C12N15/09, C07K14/475, C07K16/18, C12N15/00
CC Key Location/Qualifiers
FT source /organism='Homo sapiens (hum
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 10)

Hashimoto,S., Matsushima,K. and Suzuki,T.
Genes with human dendritic cell expression
Patent: JP 2000279181-A 276 10-OCT-2000;
                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Genes with human dendritic cell expression Patent: JP 2000279181-A 22 10-OCT-2000; SCIENCE & TECH AGENCY
                                                                                                         E39743.1 GI:18621834
JP 2000279181-A/276.
                                                                                                                                    Genes with human dendritic E39743
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1 (bases 1 to 10)

Hashimoto, S., Matsushima, K. and Suzuki, T.
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                                                                                            Homo sapiens (human)
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/db_xref="taxon:9606"
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Pred. No. 51;
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RESULT 69
AX623125
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OS Homo sapiens (hum
PN JP 2000279181-A/2
PD 10-OCT-2000
PF 01-APR-1999 JP 199
PR
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 AX623125
Sequence
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1 (bases 1 to 10)

Matsushima, K., Hashimoto, S., Kaneko, S. and Yamashita, T. Human normal liver cell expression genes Patent: JP 2001211883-A 181 07-AUG-2001; SCIENCE & TECH AGENCY
OS Homo sapiens (human)
PN JP 2001211883-A/181
PD 07-AUG-2001 JP 2000023170
PF 31-JAN-2000 JP 2000023170
PF 31-JAN-2000 JP 2000023170
PF STANDAMA, SHINICHI HASHIMOTO, SHUICHI KANEKO, TARO PI
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EH Key Location/Qualifiers
FT source 1..10
ft /organism='Homo sapiens (human)'.
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JP 2000279181-A/276
10-OCT-2000
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/mol_type="genomic DNA"
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RESULT 71
AX393189
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AX630546
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                                                 Homo sapiens
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AX630546
                                                                                                    Sequence 119 from Patent AX393189
                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                               Homo sapiens (human)
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Homo sapiens
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae,
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St Croix, B.,
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AX623125.1 GI:28451066
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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7587 from Patent WO02053774.
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Kinzler, K.W. and Vogelstein, B
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Sequence 328 from Patent
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Homo sapiens
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Sequence 214 from Patent
AX470637
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Patent: WO 0210217-A 119 07-FEB-2002;
The Johns Hopkins University (US)
Location/Qualifiers
                                                                                        Hofmann,K., Conradt,M. and Petersohn,D. Method for determining skin stress or skin Patent: WO 02053773-A 328 11-JUL-2002;
                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Sequence 916 from Patent WO02053773.
AX471339
AX471339.1 GI:22206464
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AX471385
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Method for determining skin stress or (
Patent: WO 02053773-A 916 11-JUL-2002;
HENKEL KGAA (DE)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
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Similarity 90.0%;
9; Conservation
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Means and methods for treatment evaluation
Patent: EP 1225233-A 4 24-JUL-2002;
Amsterdam Support Diagnostics B.V. (NL)
Location/Qualifiers
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Homo sapiens (human)
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AX623449
            Petersohn,D., Conradt,M. and Hofmann,K. Method for determining homeostasis of the sk Patent: WO 02053774-A 823 11-JUI-2002; Henkel Kommanditgesellschaft auf Aktien (DE)
                                                                                Homo sapiens (human)
Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Sequence 823 from Patent
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                  Petersohn,D., Conradt,M. and Hofmann,K. Method for determining homeostasis of the skin Patent: WO 02053774-A 2932 11-JUL-2002; Henkel Kommanditgesellschaft auf Aktien (DE) Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
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/mol_type="genomic DNA"
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Sequence 7314 from Patent
AX630273
AX630273.1 GI:28458311
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AX630156
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
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              Method for determining homeostasis of the skin Patent: WO 02053774-A 7911 11-JUL-2002;
                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 7911 from Patent
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AX630870.1 GI:28458910
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                         Petersohn, D., Conradt, M. and Hofmann, K.
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                                                                              Viruses; dsDNA viruses, no RNA stage; Adenoviridae; Mastadenovirus.

1 (bases 1 to 11)

Hearing, P., Schmid, S.I., Ostapchuk, P.H. and Erturk, E.
Selective regulation of adenovirus production
Patent: JP 2002506355-A 22 26 FEB-2002;
THE RESEARCH FOUNDATION OF STATE UNIVERSITY OF NEW YORK
OS Adenovirus
PN JP 2002506355-A/22
PD 26 FEB-2002
PF 15-APR-1999 JP 1999552110
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Selective regulation of adenovirus
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JP 2002506355-A/22.
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AX631203
                                                    PF 15-APR-1999 JP 1999552110
PR 15-APR-1998 US 60/081867,05-JUN-1998 US 60/08832:
PATRICK HEARING,SUSANNE I SCHMID,PHILONIENA H OSTAPCHUK,ECE
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Patent: WO 02053774-A 8245 11-JUL-2002;
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15-APR-1999
15-APR-1998
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
4 c 2 g
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Patent: EP 0939122-A 2 01-SEP-1999;
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A94680
                                                                      PATENT: WO 0018913-A 2 06-APR-2000;
ES JOHANNES HENDRIKUS VAN (NL); UNIV UTRECHT
JOHANNES CAROLUS (NL); PELFER MARK ALAN (US)
Location/Qualifiers
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/db_xref="taxon:10535"
/db_3 ~ 3 g
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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Location/Qualifiers
/organism="synthetic construct"
/mol type="genomic DNA"
/db xref="taxon:32630"
/note="mutation Fcf-responsive element"
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Sequence 2376 from Patent
AX668927
AX668927.1 GI:29291904
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AX668925.1 GI:29291902
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Biosciences Inc. (US)
Location/Qualifiers
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Biosciences Inc. (US)
Location/Qualifiers
                                                                         /note="example target 3 c 4 g
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
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Patent: WO 0138577-A 1228 31-MAY-2001,
The Johns Hopkins University (US)
                                                                                                                                                                                                                                                                                     Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 1228 from Patent
AX153313
AX153313.1 GI:14534964
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Biosciences Inc. (US)
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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AX153316
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Sequence 1230 from Patent
AX153315
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Patent: WO 0138577-A 1230 31-MAY-2001;
The Johns Hopkins University (US)
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Patent: WO 0138577-A 1229 31-MAY-2001;
The Johns Hopkins University (US)
 Human transcriptomes
                                       Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates;
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Mammalia; Eutheria; Primates;
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            Velculescu, V.E., Vogelstein, B.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/db_xref="taxon:9606"
3 c 4 g
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/mol_type="genomic DNA"
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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                                       Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
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Patent: WO 0138577-A 1233 31-MAY-2001;
The Johns Hopkins University (US)
Location/Qualifiers
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Patent: WO 0138577-A 1232 31-MAY-2001;
The Johns Hopkins University (US)
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The Johns Hopkins University (US)
Location/Qualifiers
                                                                                                                  Velculescu, V.E., Vogelstein, B. and Kinzler, K.W.
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/mol_type="genomic DNA"
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        /organism="Homo sapiens"
/mol_type="genomic DNA"
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/mol_type="genomic DNA"
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OS Homo sapiens (human)
PN JP 2001327293-A/45
PD 27-NOV-2001
PF 22-MAY-2000 JP 2000150562
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Sequence
AX153319
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Patent: WO 0138577-A 1234 31-MAY-2001;
The Johns Hopkins University (US)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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27-NOV-2001
22-MAY-2000 JP 2000150562
KOJI MATSUSHIMA, SHINICHI P
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1234 from Patent WO0138577
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                         /organism="Homo sapiens"
/mol_type="genomic DNA"
                                                                 Location/Qualifiers
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BD161262
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PD 02-UUL-2002
PF 19-DEC-2000 JP 200385816
PI SHIGENORI NAGAL, KOJI MATSUSHIMA, SHINICHI HASHIMOTO PC C12N15/09, C07K14/47, C07K16/18, C12P21/08, C12N15/00 CC Humar activated Thi and Thi cell expression genes FH Key Location/Qualifiers
FT source 1. 10
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                  1 (bases 1 to 10)

1 (bases 1 to 10)

Matsushima, K., Hashimoto, S., Kaneko, S. and Yamashita, T.

Human liver disease-expressing genes
Patent: JP 2002209591-A 757 30-JUL-2002;

JAPAN SCIENCE AND TECHNOLOGY CORP
OS Homo sapiens (human)
PN JP 2002209591-A/757
PD 30-JUL-2002
PF 19-JAN-2001 JP 2001012328
PF 19-JAN-2001 JP 2001012328
PF KOJI MATSUSHIMA, SHINICHI HASHIMOTO, SHUICHI KANEKO, TARO
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1 (bases 1 to 10)
Nagai,S., Matsushima,K. and Hashimoto,S.
Human activated Th1 and Th2 cell expression
Patent: JP 2002186482-A 84 02-JUL-2002;
JAPAN SCIENCE AND TECHNOLOGY CORP
OS Homo sapiens (human)
                                                                                                                                                                                                  unidentified.
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Human liver disease-expressing
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BD167212.1 GI:27873024
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BD161262.1 GI:27867020
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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2N15/09, C07K14/47, C07K16/18, G01N33/15, G01N33/50//C12P21/02,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism='Homo
Location/Qualifiers
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E54829
E54829.1 GI:22556312
JP 2001211883-A/181.
Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                 Genes with human dendritic cell expression Patent: JP 2000279181-A 22 10-OCT-2000; SCIENCE & TECH AGENCY
                                            E54829 10 bp Human normal liver cell expression
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (Dases 1 to 10)
Hashimoto,S., Matsushima,K. and Suzuki,T.
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Homo sapiens (human)
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C12N15/00
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JP 2000279181-A/22
10-OCT-2000
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Key Location/Qualifiers
                                                                                                                                                                                                                                                          /organism='Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="unidentified"
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/mol_type≈"genomic DNA"
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S Matsushima,K., Hashimoto,S., Kaneko,S. and Yamashita,T. Human normal liver cell expression genes
L Patent: JP 2001211883-A 181 07-AUG-2001;
SCIENCE & TECH AGENCY
OS Homo sapiens (human)
PN JP 2001211883-A/181
PD 07-AUG-2001 JP 2000023170
PF 31-JAN-2000 JP 2000023170
PI KOJI MATSUSHIMA,SHINICHI HASHIMOTO,SHUICHI KANEKO,TAR
                                                                          Sequence
AR088590
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
AR058606
          Unknown.
Unclassified.
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                                                                                                                                                                                                                                                                                                                    Unclassified.

1 (bases 1 to 10)
Chee, M., Cronin, M.T., Fodor, S.P.A., Huang, X.X., Hubbell, E.A., Lipshutz, R.J., Lobban, P.E., Morris, M.S. and Sheldon, E.L. Arrays of nucleic acid probes on biological chips Patent: US 5837832-A 183 17-NOV-1998;
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KOJI MATSUSHIMA, SHINICHI HASHIMOTO, SHUICHI KANEKO, TARO
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Thompson, J.D. Method and reagent for inhibiting P-glycoprotein (mdr-1-gene) Patent: US 5989906-A 6 23-NOV-1999;
                                                                                      The Johns Hopkins University Location/Qualifiers
                                                                                                               Human transcriptomes
Patent: WO 0138577-A 876 31-MAY-2001;
                                                                                                                                        Velculescu, V.E., Vogelstein, B. and Kinzler, K.W
                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Location/Qualifiers
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Patent: WO 0138577-A 621 31-MAY-2001;
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Patent: WO 0138577-A 1214 31-MAY-2001;
The Johns Hopkins University (US)
Location/Qualifiers
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Patent: WO 0138577-A 877 31-MAY-2001;
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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ACCESSION
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BD007922
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ACCESSION
                Query Match
Best Local
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                                                                                                                    source
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OS Homo sapiens (human)
PN JP 200106993-A/198
PD 21-MAR-2001
PF 28-APR-2000 JP 2000131079
PR
PR
FR KOJI MATSUSHIMA, SHINICHI HASHI)
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LPS activated human monocyte
BD007922
BD007922.1 GI:18636295
JP 2001069993-A/198.
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Human transcriptomes
Patent: WO 0138577-A 1215 31-MAY-2001;
Patent: WO 0138577-A University (US)
                                                                                                                                                                                                                                                                                                                                                         Matsushima, K., Hashimoto, S. and Suzuki, T. LPS activated human monocyte expressing genes Patent: JP 2001069993-A 198 21-MAR-2001;
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Homo sapiens
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1215 from Patent AX153300
                                                                                                                                                   걸걸꾼은
                                                                                                                                                                                                                                      PI KOJI MATSUSHIMA,SHINICHI HASHIMOTO,TAKUJI SUZUKI PC
C12N15/09,C07K14/47,C07K16/18,G01N33/50,G01N33/53//A61K45/00,
                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 10)
Similarity
8; Conserv
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8; Conserv
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Location/Qualifiers
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 Conservative
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                                                                                                                              /organism='Homo
                                                       /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/db_xref="taxon:9606"
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100.0%; Pred. No. 63,
Live 0; Mismatches
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                Score 8; ; Pred. No.
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SOURCE
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AUTHORS
TITLE
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BD083213
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KEYWORDS
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BD007953/c
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TITLE
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Best Local Similarity
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                                                                                                                                                                                                                                                                          ORGANISM
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                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleo Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

E 1 (bases 1 to 10)

E Matsushima, K., Hashimoto, S., Suzuki, T. and Nagai, S.
Human matured/activated dendritic cell expression genes
L Patent: JP 200137793-A 134 27-NOV-2001;
L DAPAN SCIENCE AND TECHNOLOGY CORP
OS Homo sapiens (human)
PN JP 200137793-A/134
PD 27-NOV-2001
PF 22-MAY-2000 JP 2000150562
PF 22-MAY-2000 JP 200150562
PF 22-MAY-2000 JP 200150562
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PC A61P3
CC
FH Key
FT sourc
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(bases 1 to 10)

Matsushima, K., Hashimoto, S. and Suzuki, T.

LPS activated human monocyte expressing genes
Patent: JP 200106993-A 229 21-MAR-2001;

JAPAN SCIENCE AND TECHNOLOGY CORP

OS Homo sapiens (human)

PN JP 2001069993-A/229
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JP 2001327293-A/134.
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C12N15/09, C07K14/47, C07K16/18, G01N33/50, G01N33/53//A61K45/00,
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LPS activated human monocyte
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                                                      NAGAI
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                                   C12N15/09, C07K14/47, C07K16/18//C12P21/02, C12P21/08, C12N15/00
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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Pred. No. 63;
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Location/Qualifiers
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                                                                    SUZUKI, SHIGENORI
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REFERENCE
AUTHORS
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ACCESSION
VERSION
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BD161348
LOCUS
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ORGANISM
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SOURCE
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Best Local S
Matches 8
                          AUTHORS
TITLE
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nes 8; Conserv
         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; 1 (bases 1 to 10)
Nagai, S., Matsushima, K. and Hashimoto, S.
Human activated Th1 and Th2 cell expression genes
Patent: JP 2002186482-A 170 02-JUL-2002;
                                                                                                                     Human activated Th1 and BD161348 BD161348.1 GI:27867106 JP 2002186482-A/170.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E Mammalla; Eutheria; Primates; Catarrhini; Hominidae; 1 (Dases 1 to 10)

Matsushima, K., Hashimoto, S., Suzuki, T. and Nagai, S. Human matured/activated dendritic cell expression ger Patent: JP 2001377293 A 229 27-NOV-2001; JAPAN SCIENCE AND TECHNOLOGY CORP
                                                                                            Homo sapiens
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JP 2001327293-A/229.
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JAPAN SCIENCE AND
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KOJI MATSUSHIMA,SHINICHI HASHIMOTO,TAKUJI SUZUKI,SHIGENORI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens (human)
JP 2001327293-A/229
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                   /organism="Homo sapiens"
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/db_xref="taxon:9606"
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PC C12N;
PC C12N;
PC C12N;
CC Human
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PN JP 2002186482-A/170
PD 02-JUL-2002
PD 19-DEC-2000 JP 2000385816
PF 19-DEC-2000 JP 2000385816
PT 
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Matsushima, K., Hashimoto, S., Kaneko, S. ar
Human liver disease-expressing genes
Patent: JP 2002209591-A 117 30-JUL-2002;
JAPAN SCIENCE AND TECHNOLOGY CORP
OS Homo Sapiens (human)
PN JP 2002209591-A/117
BD167054
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Key Location/Qualifiers
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19-JAN-2001 JP 2001012328
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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o_xref="taxon:32644"
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Homo sapiens (human)
Homo sapiens
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Human normal liver cell expression genes
Patent: JP 20012111883-A 67 07-AUG-2001;
SCIENCE & TECH AGENCY
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E54715.1 GI:22556198
JP 2001211883-A/67.
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JP 2002209591-A/599
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JP 2001211883-A/67
07-AUG-2001
31-JAN-2000 JP 2000023170
KOJI MATSUSHIMA,SHINIÇHI HA
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KOJI MATSUSHIMA,SHINICHI HASHIMOTO,SHUICHI KANEKO,TARO
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                                                                              Location/Qualifiers.
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/db_xref="taxon:32644"
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            db_xref="taxon:9606"
                                organism="Homo sapiens"
/mol_type="genomic DNA"
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Catarrhini; Hominidae
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AX511277
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AX482038
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Sequence 15 from Patent EP1225233.
AX482038
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Means and methods for treatment evaluation
Patent: WO 02059558-A 15 01-AUG-2002;
Amsterdam Support Diagnostics B.V. (NL)
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Means and methods for treatment evaluation
Patent: EP 1225233-A 15 24-JUL-2002;
Amsterdam Support Diagnostics B.V. (NL)
Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
Homo sapiens (human)
Homo sapiens
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/mol_type="genomic DNA"
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Blledge,S.J. and Sanchez,Y.
Balledge,S.J. and Sanchez,Y.
Mammalian checkpoint genes and proteins
Patent: US 6307015-A 6 23-CCT-2001;
Location/Qualifiers
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Elledge, S.J. and Sanchez, Y.
Mammalian checkpoint genes and proteins
Patent: US 6218109-A 6 17-APR-2001;
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Monia, B.P. and Xu, X.S.
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MCKAY, R., Dean, N., Monia, B.P., Nero, P.S. and Gaarde, W.A.

Antisense oligonucleotide compositions and methods for the

modulation of UNK proteins

Patent: US 6133246-A 31 17-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Methods of modulating tumor necrosis factor expression of cell adhesion molecules Patent: US 6114517-A 7 05-SEP-2000;
Location/Qualifiers
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patent US
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Pred. No.
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Sequence 2 from Patent WO0134792.
AXI64692
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Patent: US 6133246-A 42.17-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         synthetic construct
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Patent: WO 0134792-A 2 17-MAY-2001;
THE SECRETARY OF THE DEPARTMENT OF HEALTH
                                                                                                                                         artificial sequences.
                                                                                                                                                      synthetic construct synthetic construct
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
/db_xref="taxon:32630" 2 t
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AX355435.1 GI:18620103
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10; Conserv
                                                                                                                         Brattler,R.L. Inhibition of angiogenesis by nucleic Inhibition of angiogenesis by nucleic Patent: WO 02053141-A 311 11-JUL-2002; Coley Pharmaceutical Group, Inc. (US)
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
/note="Synthetic Sequence"
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/db xref="taxon:32630"
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Patent: JP 2001514905-A 31 18-SEP-2001;
ISIS PHARMACEUTICALS INC
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JP 2001514905-A/42.
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Artificial Sequence
JP 2001514905-A/42
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nilarity 71.4%;
Conservative . (
18-SEP-2001
07-AUG-1998 JP 2000509875
13-AUG-1997 US 08/910629
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ROBERT MCKAY,NICHOLAS DEAN,BRETT P MONIA,PAMELA SCOTT
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/db_xref="taxon:32630"
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composition and modulation method
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AX668699
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NERO,WILLIAM A GAARDE
PC C1201/68,A61K31/7088,A61K48/00,A61P35/00,C12N15/09,C12P19/34,
PC C12N15/00
CC control sequence (sense)
FH Key Location/Qualifiers
FT source 1.20
FT source /organism='Artificial Sequence'.
Liu,Q.
Position dependent recognition fingers
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1 (bases 1 to 9)
Mauro, V.P., Edelman, G.M., Chappell, G.M.,
Krushel, L. and Zhou, W.
Synthetic internal ribosome entry sites (
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                                                                                                        artificial sequences
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8; Conserv
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
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Sangamo
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Sequence 11 from Patent WO0160997.
AX224404
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                                                      Male tissue-preferred regulatory region Patent: WO 0160997-A 11 23-AUG-2001; PIONEER HI-BRED INTERNATIONAL, INC. (US)
                                                                                                                       Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; PACCAD clade; Panicoideae; Andropogoneae; Zea.
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Biosciences Inc. (US)
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/db_xref="taxon:32630"
/note="example target DNA"
2 C 6 g 0 1
/organism="Zea mays"
/mol_type="genomic DNA"
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/mol_type="genomic DNA"
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Patent: WO 0138577-A 1215 31-MAY-2001;
The Johns Hopkins University (US)
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Patent: WO 0138577-A 1214 31-MAY-2001;
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/mol_type="genomic DNA"
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LPS activated human monocyte expressing genes
Patent: JP 200106993-A 229 21-MAR-2001;
JAPAN SCIENCE AND TECHNOLOGY CORP
OS Homo sapiens (human)
PN JP 200106993-A/229
PD 21-MAR-2001 JP 2000131079
PF 28-APR-2000 JP 2000131079
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C12N15/09, C07K14/47, C07K16/18, G01N33/50, G01N33/53//A61K45/00,
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Method for determining homeostasis of the skin
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AX129247
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Sequence 13 from Patent W09947656.
AX017438
AX017438.1 GI:10042250
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Ribozyme therapy for the treatment of proliferative skin and
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Sequence 465 from Patent WO0130362.
AX129247
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                                                                                                                                                                                                                                                                                                                         Patent: WO 0130362-A 465 03-MAY-2001;
                                                                                                                                                                                                                                                                                                                                      diseases
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                     /note="Cdk4 ribozyme binding site"
6 c 7 g 5 t
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                                                                                                                                                                                                                                             organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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AX668947/c
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AX538878/c
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M synthetic construct
Artificial sequences.
1 (bases 1 to 9)
1 Isalan,M. and Choo,Y.
Nucleic acid binding proteins
Patent: WO 9947656-A 13 23-SEP-1999;
MEDICAL RES COUNCIL (GB); ISALAN MARK
Location/Qualifiers
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                                                                                                         Sequence 23
AX668947
AX668947.1
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Mini-adenoviral vector system for vaccination
Patent: WO 0208436-A 12 31-JAN-2002;
Genstar Therapeutics Corporation (US)
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Sequence 12 from Patent WO0208436.
XX538878
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 Patent:
            fingers
                      Position dependent recognition of gnn nucleotide triplets by zinc
                                    Liu,
                                                           artificial sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Viruses; dsDNA viruses, no RNA stage; Adenoviridae;
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WO 0242459-A 2396 30-MAY-2002;
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/mol_type="genomic DNA"
/db_xref_"taxon:32630"
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                                                                                                                                                                                                                                                                                                                    /note="A repeat VI"
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/mol_type="genomic_DNA"
/db_xref="taxon:10535"
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OS Artifi
PN JP 2000
PD 05-MAR
PF 17-MAR
PR 17-MAR
03-APR-1998
PI YEN CH
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Choo, Y. and Isalan, M.
Nucleic acid binding proteins
Patent: JP 2002506640-A 4 05-N
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synthetic construct
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JP 2002506640-A/4.
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Nucleic acid binding proteins.
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Sangamo Biosciences Inc. (US)
Location/Qualifiers
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Sequence 2397 from Patent WO0242459
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                 -APR-1998 GB
                17-MAR-1999 JP 2000536839
17-MAR-1998 GB 9805576.7,31-MAR-1998
PR-1998 GB 9807246.5
                                                    Artificial Sequence
JP 2002506640-A/4
05-MAR-2002
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Location/Qualifiers
     CHOO, MARK ISALAN
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AX471339/c
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AX471339
AX471339.1 GI:22206464
                                                   Method for determining homeostasis of the sh
Patent: WO 02053774-A 823 11-JUL-2002;
Henkel Kommanditgesellschaft auf Aktien (DE)
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                                                                                          Petersohn,D., Conradt,M. and Hofmann,K.
                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
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/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/organism="Homo sapiens'
/mol_type="genomic DNA"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Patent: WO 0138577-A 876 31-MAY-2001;
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/mol_type="genomic DNA"
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Location/Qualifiers
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Patent: WO 0138577-A 877 31-MAY-2001;
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THORS Tomizuka, K., Yoshida, H., Ishida, I. and Kuroiwa, Y.

Non-human animal having modified foreign chromosomal or the construction of the 
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PN JP 2001231403-A/6
PD 28-AUG-2001
PF 18-FEB-2000 JP 2000042074
PI KAZUWA TOWIZUKA, HITOSHI YOSHIDA, ISAO ISHIDA, YOSHIMI KUROIWA PC A01K67/027, C12N5/10, C12N15/09//(C12N5/10, C12R1:91),
PC C12N5/00, C12N15/00 (C12N6/10, C12N6/10, C12R1:91)
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Non-human animal having modified foreign chromosomal or slice
Patent: JP 2001231403-A 6 28-AUG-2001;
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T KAZUWA TOWIZUKA, HITOSHI YOSHIDA, ISAO ISHIDA, YOSHIMI KUROIWA PC A01K67/027, C12N5/10, C12N15/09/ (C12N5/10, C12R1:91), (C12N15/09, PC C12R1:91)

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Description of Artificial Sequence: SrfI linker FH Key

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Antisense oligonucleotide; Jun N-terminal kinase; JNK; hybridise; JNK1; JNK2; JNK3; cell cycle progression; phosphorylation; tumour; probe; hyperproliferative disease; human; ss.
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Matches 20
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                WPI; 1999-181060/15
                                                                                                                                            07-AUG-1998;
                                                                                                                                                                                                                                                                                   Antisense oligonucleotide; Jun N-terminal kinase; JNK; hybridise; JNK1; JNK2; JNK3; cell cycle progression; phosphorylation; tumour; probe; hyperproliferative disease; human; ss.
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                                                                               (ISIS-)
                                                                                                              13-AUG-1997;
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RESULT 1

New antisense oligonucleotides that detect and modulate

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense; gene therapy; JNK2 protein; apoptosis; cancer; cellular hyperproliferation; Alzheimer's; Parkinson's disease; amylotrophic lateral sclerosis; retinitis; pigmentosa; epilepsy; myocardial infarction; stroke; obstructive jaundice; polycystic kidney; diabetes; Jun N-terminal kinase; ss.
The present invention relates to antisense oligonucleotides (AACG2844-C63000, AAA96093-A96099 and AAA07993) that hybridise specifically to a nucleotide encoding a Jun N-terminal kinase protein, resulting in decrease of JNK2 expression and leading (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to antisense oligonucleotides that detect and modulate the expression of Jun N-terminal kinase (JNK) proteins. The oligonucleotides specifically hybridize to a nucleic acid encoding JNK1, JNK2 or JNK3 protein, and which modulate expression of these
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                                                                                                                                          Novel methods for reducing apoptosis comprising contacting cells with antisense oligonucleotides, useful for treating apoptotic disorders,
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oligonucleotide. The oligonucleotides of the present invention are a for treating diseases or conditions with reduced apoptosis, e.g. can and cellular hyperproliferation. The oligonucleotides may also be us increase the stimulation of apoptotic proteins, e.g. for treating Alzheimer's or Parkinson's disease, amylotrophic lateral sclerosis, retinitis, pigmentosa, epilepsy, myocardial infarction, stroke, obstructive jaundice, polycystic kidney and diabetes. The present sequence may have a phosphorothioate backbone.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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WO200059549-A1 sapiens.

04-APR-2000; 2000WO-US08880

12-OCT-2000

07-APR-1999; 99US-0287796

(ISIS-) ISIS PHARM INC.

Dean NM. Monia ₽P, Nero PS, Gaarde

2000-638427/61.

Novel methods for reducing antisense oligonucleotides, apoptosis , useful fo is comprising for treating contacting cells with apoptotic disorders,

Example 4; Page 135; 160pp; English

induction of apoptosis. The present sequence is one such antisense oligonucleotide. The oligonucleotides of the present invention are useful for treating diseases or conditions with reduced apoptosis, e.g. cancer and cellular hyperproliferation. The oligonucleotides may also be used to increase the stimulation of apoptotic proteins, e.g. for treating Alzheimer's or Parkinson's disease, amylotrophic lateral sclerosis. The present invention relates to antisense oligonucleotides (AAC62844-C63000, AAA96093-A96099 and AAA07993) that hybridise specifically to a nucleotide encoding a Jun N-terminal kinase (JN protein, resulting in decrease of JNK2 expression and leading to retinitis, pigmentosa, epilepsy, myocardial infarction obstructive jaundice, polycystic kidney and diabetes. sequence may have a phosphorothioate backbone. infarction, stroke (JNK2

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RESULT 5
ADA448651
ID ABA44
XX AAA4
XX AAA44
XX AAA44
XX AAA4
XX AAA44
XX AAA4
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Best Local
                       signalling molecule. In the method Ha-ras, c-raf and c-Jun N-terminal
                                                                                                                                                                         Modulating cell adhesion molecule expression for treating immune or inflammatory diseases involves treating cell with specific inhibitor of Tumour Necrosis Factor alpha signalling molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antisense; E-selectin; TNF alpha; cell adhesion; tumour necrosis factor alpha; phosphorothioate; methoxyethoxy; sepsis; rheumatoid arthritis; inflammatory; immune disease; inflammatory bowel disease; allograft rejection; cancer; antibacterial; diabetes; Grave's disease; allograft rejection; cancer; antibacterial; immunosuppressive; antipsoriatic; antidiabetic; antithyroid; cytostatic; dermatological; antiallergic; Ha-ras; c-raf;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                     Claim
                                                                                                                                                                                                                                                                           WPI; 2000-423367/36.
                                                                                                                                                                                                                                                                                                                     Monia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense oligonucleotide ISIS no.15354 to human JNK2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAA48651;
                                                                 A novel method for modulating cell involves antisense inhibition of a
                                                                                                                                                                                                                                                                                                                                                                  (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                               10-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-JUN-2000
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                                                                                                                                     36;
                                                                                                                                                                                                                                                                                                                     ₽,
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                                                                                                                                   Page 46; 100pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                                                                                                     XS;
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16..20
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/note= "All bases are
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/mod_base= OTHER
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/note= "All bases are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                *tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
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                                                                                                                                   English.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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..
                  adhesion molecule expression
tumour necrosis factor (TNP) alp
TNF alpha signalling molecules
kinase (JNK) 2 were inhibited by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           internucleotide linkage"
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RESULT 6
AAH23754
ID AAH2
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XX AAH2
XX AAH2
XX JNK1
XX JNK1
XX JNK1
XX Synt
XX Synt
XX Synt
XX Synt
XX HO2(
XX Synt
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Best Local :
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                     The present invention relates to the use of J oligonucleotides for treating cancer and for mimic or augment the effect of JNK antisense for cancer. The present sequence is one such
                                                                                                                                                      Claim 1;
                                                                                                                                                                                                                                 Use of Jun Kinase antisense mRNA for treating cancer by administering vector comprising promoter operably linked to DNA sequence that encod
                                                                                                                                                                                                                                                                                                               WPI;
                                                                                                                                                                                                                                                                                                                                                                Potapova 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-NOV-2000; 2000WO-US30869
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rejection and cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                   (USSH ) US DEPT HEALTH & HUMAN SERVICES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RK; jun kinase; antisense; cytostatic;
-O-methoxyethyl oligonucleotide; MOE;
                                                                                                                                                                                                         antisense mRNA to patient
                                                                                                                                                                                                                                                                                                               2001-335925/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antisense
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                                                                                                                                                      41; 75pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          chimeric antisense oligonucleotide containing MOE/phoshopdiester residues flanking a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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Pred. No. 1.1;
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                                                                                                                                                                                                            with
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or screening compounds
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th JNK antisense
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Matches
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                                                                                                            immune response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells.

Note: the present sequence may have a phosphorothioate backbone.
                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                               Vaccinating against tumors, infectious diseases, allergies and using immunostimulatory Py-rich and TG nucleic acids -
                                                                                                                                                                                                                                                                                                                                                                                                                                      Krieg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-SEP-1999; 99US-0156113.
27-SEP-1999; 99US-0156135.
23-AUG-2000; 2000US-0227436.
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                                                                                                                                                                                                                                                                         response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an
                                                                                                                                                                                                                                                                                                                                  Claim 101; Page 44; 338pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vaccine; cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma; infectious disease; allergy; immune deficiency; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF99183;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAF99183 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunostimulatory nucleic acid #299.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (COLE-)
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                                          Similarity
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GTCCGGGCCAGGCCAAAGTC 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Schetter C,
                            Conservative
                                                                                  BP; 4 A; 7 C; 7 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                       invention relates to a method for stimulating an immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                           Matches
                                                                                                                                                      Query Match
Best-Local
                                                                                                                                                                                                                                                 Sequence 20 BP; 4 A; 7 C; 7 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                     a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic acid of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; obler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Inhibiting angiogenesis in a subject, involves administering one antiangiogenic nucleic acid molecule to the subject -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (COLE-) COLEY PHARM GROUP INC
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                                                                                                                                                         Similarity
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Pred. No. 1
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RESULT 9 ABL39057 ID ABL3

ABL39057 standard; DNA; 20

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RESULT 10
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Matches 20
                                                                                                                                                                                                                                                                                                         treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, breast cancer, cervical cancer, colon and rectum cancer, connective tissue cancer, ossophageal cancer, eye cancer, kidney cancer, larynx cancer, leukaemia, liver cancer, lung cancer, Hodgkin's lymphoma, mon-Hodgkin's lymphoma, melanoma, myeloma, oral cavity cancer, ovaria cancer, promotate cancer, rhabdomyosarcoma, skin cancer, stomach cancer, testicular cancer, and uterine cancer. The present sequence is an immunostimulatory oligonucleotide described in
             08-DEC-1998
                                           AAV60744.;
                                                                                                                                                                                                                                                                 Sequence 20 BP; 4
                                                                                                                                                                                                                                                                                              cancer, stomach cancer, testicular capresent sequence is an immunostimulat the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to methods for treating or preventing cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies. The methods are useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies to a subject having or at risk of
                                                                     AAV60744 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 212; 312pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-154611/20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antibody-induced cell lysis; cancer; immunostimulatory; CD20; angiogenesis; metastasis; cytostatic; phosphorothioate backbone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (IOWA ) UNIV IOWA RES FOUND
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                                                                                                                                                                                                         20;
                                                                                                                                                                                                                     Similarity
                                                                                                                                             GTCCGGGCCAGGCCAAAGTC 20
                                                                                                                                                                          GTCCGGGCCAGGCCAAAGTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hartmann G;
                                                                                                                                                                                                         Conservative
            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mod
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        *tag=
                                                                                                                                                                                                                                                                 A; 7 C; 7 G; 2 T; 0 other;
                                                                     DNA;
                                                                                                                                                                                                                     100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          base= OTHER
                                                                                                                                                                                                                                   100.0%;
                                                                     18
                                                                     ВÞ
                                                                                                                                                                                                        0
                                                                                                                                                                                                                     Score 20;
Pred. No.
                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                        0
                                                                                                                                                                                                                                  Length 20;
                                                                                                                                                                                                        Indels
                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                        ovarian
                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               88
                                                                                                                                                                                                      0
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Screening for agents that effect cell cycle regulatory proteins using a cell line that expresses a reporter gene in response to regulation through phosphorylation by a cyclin/CDK system
                                                                                                                                                                                                                                                                   WPI; 1998-251302/22
                                                                                                                                                                                                                                                                                                                          27-NOV-1996;
16-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                          23-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                           WO9816660-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                          cyclin kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Primer
                                                                                                                                                                                                                                                                                                                                                       16-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                           transcription
                                                                                                                                                                                                                                                                                                       (BITT-) BITTECH INC
                                                                                                                                                                                                                                                                                                                                                                                                                                        primer; amplification; yeast; UAS; upstream activating sequence; upstream activating sequence; UAS; nscription terminator; cell cycle; Upstream Activation Sequence; UAS; moter; phosphorylation; cyclin; dependent kinase; CDK; vector; lin kinase inhibitor; CKI; growth; wound healing; cancer therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        #2 for human
                                                                                                                                                                                                                                                                                                                        96US-0029127.
                                                                                                                                                                                                                                                                                                                                                       97WO-US18608
                                                                                                                                                                                                                                                                                                                                     96US-0031968
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CDK4 codons 1-163.
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CC linked to codons 154-302 of the yeast PHO85 gene. The fusion protein is Cused to screen for compounds that affect mammalian cell cycle regulatory proteins. The method comprises administering a compound to a cell line, compound to a cell compound to a cell compound to a cell compound that compound the construct is an effector gene compounding a gene product to permit normal cyclin/CDK regulation of the compound affects the normal regulation. The method can be used to identify inhibitors and activators of cyclins, cDKs, cyclin/CDK complexes, cyclin kinase compound that cell cycle regulatory proteins, especially inhibitors and activators of cyclins, CDKs, cyclin/CDK complexes, cyclin kinase can be used for stimulating growth of cells (as in wound healing), or can be used for stimulating growth and division (as in cancer therapy). Primers AAV60743-V60745 were used to PCR amplify codons 1-163 of the human cyclin-dependent kinase 4 (hCDK4). The amplified product was used to generate a fusion protein comprising part of the hCDK4 sequence linked to codons 154-302 of the yeast PHO85 gene. The fusion protein is

Example 4; Page 75; 93pp; English.

Sequence 18 BP; 4 A; 6 C; 5 G; 3 T; 0 other;

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밁
            Ś
                           Query Match
Best Local S
Matches 16
N
                                  Similarity
            CGGGCCAGGCCAAAGTC
                            Conservative
                                  77.0%;
            20
                            0
                       Pred. No. 10,
0; Mismatches
                                         Score 15.4;
                                        DB 1;
                           1;
                                        Length 18;
                           0
                          Gaps
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RESULT 11
AAA82879/c
ID AAA828
XX
AAA828
AC AAA828
XX
DT 04-DEC
XX
DE Cdk4 r
XX
XW
Ribozy
Ribozyme; hairpin; hammerhead; gene therapy; vasotropic;
                        cdk4 ribozyme binding site #60
                                                04-DEC-2000
                                                                        AAA82879
                                                                                                AAA82879 standard;
                                                (first entry)
                                                                                                DNA;
                                                                                                 19
                                                                                                ВP
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RESULT 12
AAH58041/c
    엉
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                              matrix metalloproteinase; growth factor; reductase; scarring; cytostatic; antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide; antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma; basal cell carcinoma; seborrheic wart; vitreoretinopathy; scar;
                                                                                                                                                                                                recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          designed to cleave RNA encoding a cyclin or cell-cycle dependent other than cell-cycle dependent kinases CDKI, PCNA and Cyclin B1. Representative examples of riboxyme cognition sites are given in AAA82415 to AAA86787. The riboxyme of the invention is useful for inhibiting restenosis by introduction of the riboxyme into cells. The riboxyme is resistant to endonuclease activity and hence is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI;
                            03-MAY-2001
                                                                                                                                                                                                                                                                    Cell-cycle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19
26-OCT-2000; 2000WO-US29500.
                                                                                 Synthetic
                                                                                                                       sickle cell
                                                                                                                                                                                                                                        Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme;
                                                                                                                                                                                                                                                                                                                         AAH58041;
                                                                                                                                                                                                                                                                                                                                                 AAH58041 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 53; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New hairpin and hammerhead ribozyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tritz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-JUN-2000
                                                    WO200130362-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (IMMU-) IMMUSOL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200032765-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               restenosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            encoding a cyclin or cell-cycle and Cyclin B1
                                                                                                                                                                                                                                                                                                                                                                                                                  19
                                                                                                                                                                                                                                                                                                                                                                                                                                           9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                         GGCCAGGCCAAAGTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ij
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Welch PJ,
                                                                                                                                                                                                                                                                dependent kinase cdk4 ribozyme binding site SEQ ID NO:465
                                                                                                                      retinopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             invention relates to a hairpin or hammerhead ribozyme, cleave RNA encoding a cyclin or cell-cycle dependent cell-cycle dependent kinases CDKI, PCNA and Cyclin B1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 1 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               restenosis treatment.
                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98US-0110954.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99WO-US28772
                                                                                                                                                                                                                                                                                                                                                                                                                                                              75.0%; or
100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6 C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Barber JR,
                                                                                                                                                                                                                                                                                                                                                 19
                                                                                                                       S
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                                                                                                                                                                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      G
5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Robbins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         dependent kinase other than CDK1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for inhibiting restenosis, cleaves
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  clin B1.
given in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             kinase
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RESULT 13
AAX73494/c
ID AAX734
XX AAX734
XX AOX94
AC AAX734
XX Wouse
XX Wouse
XX Wouse
XX Wouse
XX Wouse
XX Woscul
KW Fik-1;
KW Fis-1;
KW fims-1;
KW foetal
XX Wo9715
XX W09715
XX W09715
XX YO9715
XX PP 25-OCT
XX PR 11-JAN
PR 11-JAN
PR 11-JAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      dependent kinase, growth factor or a reductase, or administering a culleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, cophthalmological, vulnerary, keratolytic and virucide activities, and cophthalmological, cytostatic antiseborrheic, antidiabetic, antisickling, cophthalmological, vulnerary, keratolytic and virucide activities, and colleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, acopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing scarring such as keloid, adhesion and hypertrophic or hypertrophic burn scar. AAH57577 to AAH62099 represent sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                    Vascular endothelial growth factor receptor; VBGF receptor; flt-1; flk-1; KDR; hammerhad ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX73494
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19 BP; 1 A; 6 C; 7 G; 5 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Treating proliferative skin or eye diseases and scarring, using ribozymes that cleave RNA encoding cytokines involved in inflammat matrix metalloproteinases, growth factors and cell-cycle dependent
                                                          25-OCT-1996;
                                                                                                  01-MAY-1997
                                                                                                                                       WO9715662-A2
                                                                                                                                                                                                                    foetal liver kinase 1; ss.
                                                                                                                                                                                                                                                                                                                                     Mouse flk-1 VEGF receptor hairpin ribozyme substrate
                                                                                                                                                                                                                                                                                                                                                                             28-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   skin or eye disease and scarring. The method involves admini ribozyme (I) which cleaves RNA encoding a cytokine involved inflammation, matrix metalloproteinase (MMP), cyclin, cell-cinflammation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 105; 408pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2001-300427/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6 GGCCAGGCCAAAGTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ĭ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard; RNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tritz R
                                                            96WO-US17480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               9908-0161532.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  75.0%;
100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  describes a method for treating nd scarring. The method involves
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                       #41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cell-cycle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                a proliferative administering a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         inflammation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
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0,

11-JAN-1996; 26-OCT-1995;

96US-0584040 95US-0005974

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RESULT 14
AAX71472/c
ID AAX714
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local.S
Matches 16
Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage;
tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease;
fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. AAX67275 to AAX75752 represent specific examples of nucleic acid molecules from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour the flk-1) (e.g. tum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                           11-JAN-1996;
26-OCT-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim
                                                                                                                                                                                                                                                                                                                                  25-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vascular endothelial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAX71472 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1997-259017/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (CHIR )
                                                                                                                                        Escobedo J,
                                                                                                                                                                                                                                                                                                                                                                                                                             WO9715662-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                         (RIBO-)
                                                                                                                                                                                                                  (CHIR )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   KDR VEGF receptor hammerhead ribozyme substrate #484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4; Page 152; 218pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       w
                                                                                                                                                                                                                  CHIRON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CHIRON CORP.
RIBOZYME PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                       RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CCGGGCCAGGCCAAAGTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCGGGCCAAGCCGAAGTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 1 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                        McSwiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     kinase 1; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                           96US-0584040.
95US-0005974.
                                                                                                                                                                                                                                                                                                                                  96WO-US17480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             74.0%;
88.9%;
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The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or mor receptors of vascular endothelial growth factor (VBGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flk-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                       The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (VEGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase if (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase if (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase if (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase if (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase if (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase if (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase if (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase if (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase if (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase if (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase if (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase if (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase if (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase if (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase (flk-1) (e.g. tumour processors (KDR) and/or foetal liver kinase (flk-1) (e.g. tumour processors (flk-1) (e.g. tumour processor
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RESULT 17
AAD03983
ID
AAD03983
XX
AC
AAD03
AAD03
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AAD03
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AAD03
AC
AAD03
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AAD03
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AAD03
RESULT 18
AAS18424
ID AAS18
XX
AC AAS18
XX
AC AAS18
XX
DT 12-MA
XX
DE Degen
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence is a degenerate PCR primer used for amplifying the human cell cycle checkpoint protein, hchkl DNA. The cell cycle checkpoint protein, hchkl DNA. The cell cycle checkpoints are regulatory pathways that control the order and timing of cell cycle transitions, and ensure that critical events such as DNA replication and chromosome segregation are completed with high fidelity. The chkl protein controls cell cycle in response to DNA damage. It functions as kinase and phosphorylates the key regulators of Cdk tyrosine phosphorylation. The checkpoint gene sequences are used as probes for a portion of the chromosome associated with tumours and other malignancies, as well as growth and/or development deficiencies. The chkl proteins are called for generating specific antibodies and for inhibiting growth of
                             Degenerate PCR primer #2 used
                                                                                                                                                                                                    AAS18424 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New Chkl proteins and gene sequences encoding the proteins useful probes for a portion of the chromosome associated with tumors and malignancies, growth and/or development deficiencies -
                                                                                       12-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Elledge SJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US6218109-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA damage; kinase; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; cell cycle checkpoint; chkl; tumour; malignancy; cell growth inhibitor; development deficiency; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human cell cycle checkpoint protein, hchkl DNA amplifying PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAD03853 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (BAYU ) BAYLOR COLLEGE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2001-289827/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15
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                                                                                                                                                                                                                                                                                                                                                                                                   σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                          14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                 GGACAGGCCAAAGTC 17
                                                                                                                                                                                                                                                                                                                                                                                                   GGCCAGGCCAAAGTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCGGGCAAGGCCAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Column 24; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 5 A; 4 C; 5 G; 3 T; 0 other;
                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sanchez
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97US-0924183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97US-0924183
                                                                                                                                                                                                    DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     67.0%;
93.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                   20
                                                                                                                                                                                                    ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 13.4;
Pred. No. 25;
                             to amplify DNA encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                human chk1
                                                                                                                                                                                                                                                                                                                                                                                                                                                          o
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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RESULT 19
ABX16354
ID ABX16
XX ABX16
XX O8-AP
DT 08-AP
DX Human
XX Human
XX Human
XX Homo
XX US200
XX US200
XX 12-DE
XX 12-DE
XX 12-JP
XX (ELLE
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Best Local S
Matches 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              checkpoint (chk1) proteins and the nucleic acid sequences encoding them. Human chk1 (hchk1) maps to chromosome 11q24. Chk1 is involved in cellular responses to DNA damage, in the cell cycle checkpoint pathway. The protein is useful for generating specific antibodies and for inhibiting the growth of cells. The nucleotide sequence encoding the protein may be used as a probe for a portion of the chromosome associated with tumours and other malignancies, as well as growth and/or development deficiencies. The present sequence represents a degenerate PCR primer used to amplify DNA encoding human chk1 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New mammalian checkpoint protein and gene, for generating specific antibodies or for inhibiting the growth of cells, and for use as a probe for a portion of a chromosome associated with tumours or malignancies -
                                                                                                               US2002156247-A1
                                                                                                                                          Homo sapiens
                                                                                                                                                                                    Human; checkpoint;
                                                                                                                                                                                                             Human checkpoint gene Chk1 PCR primer #2
                                                                                                                                                                                                                                             08-APR-2003
                                                                                                                                                                                                                                                                                                    ABX16354 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; checkpoint protein; hchk1; DNA damage; chromosome 11q24; cell cycle checkpoint pathway; inhibition of cell growth; tumour; malignancy; growth deficiency; development deficiency; PCR primer; ss.
                          12-JAN-2000; 2000US-0488364
                                                       12-DEC-2001; 2001US-0020038.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Column 24; 39pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to the isolation of human and mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-JAN-2000; 2000US-0488364.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23-OCT-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US6307015-B1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
 (ELLE/) ELLEDGE S J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (BAYU ) BAYLOR COLLEGE MEDICINE
                                                                                                                                                                     primer;
                                                                                                                                                                                                                                                                                                                                                                          w
                                                                                                                                                                                                                                                                                                                                                                                                                                  14;
                                                                                                                                                                                                                                                                                                                                                                                                     6 GGCCAGGCCAAAGTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                          GGACAGGCCAAAGTC
                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 5
                                                                                                                                                                        88
                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sanchez
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97US-0924183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; 4 C; 5
                                                                                                                                                                                    chk1; anti-Chkl antibody; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                             67.0%;
93.3%;
                                                                                                                                                                                                                                                                                                                                                                          17
                                                                                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 13.4; Di
Pred. No. 25;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             BB
                                                                                                                                                                                                                                                                                                                                                                                                                                                            1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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RESULT 20
AAX74910/c
ID AAX749
XX AAX749
XX AOX749
XX Wouse
XX Wouse
KW Vascul
KW fik-1;
KW fims-1;
KW foetal
XX W09715
XX W09715
XX W09715
XX RH CHIR
PF 25-OCT
XX CHIR
PR 11-JAN
PR 26-OCT
XX RH
PR 11-JAN
PR 26-OCT
XX RH
PR 11-JAN
PR 26-OCT
XX CHIR
PA (RIBO-
XX RIBO-
XX Nuclei
PT Nuclei
PT Nuclei
PT Nuclei
PT Nuclei
PT PROATI
XX PSOCTIA
XX Claim
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Best Local 9
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                                                                               Nucleic acid molecule modulating VEGF receptor(s) gene expression mRNA stability - useful for treating e.g. tumour angiogenesis, psoriasis, rheumatoid arthritis, etc., in a human patient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vascular endothelial growth factor receptor; VEGF receptor; flt-1; flk-1; KDR; hammerhead ribozyme; hairpin ribozyme; cleavage; tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   binding to an antigenic determinant on the proteins end sequence comprising 476 (3 sequences), 479, 496 or 513 A new method is used to produce the antibody, which is
                                                    Claim 4;
                                                                                                                                                         WPI; 1997-259017/23.
                                                                                                                                                                                        Escobedo J,
                                                                                                                                                                                                                                                                               11-JAN-1996;
26-OCT-1995;
                                                                                                                                                                                                                                                                                                                                  25-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                    01-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                     WO9715662-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mouse flt-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAX74910
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX74910 standard; RNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 5 A; 4 C; 5 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  detecting a Chkl protein that is associated with a tumour. This sequence represents a PCR primer used to isolate DNA encoding the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention describes an anti-Chkl antibody capable of specifically
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New anti-Chkl antibody, that may be a monoclonal or polyclonal antibody, useful for detecting a Chkl protein that is associat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-182651/18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          foetal liver kinase 1; ss.
                                                                                                                                                                                                                            (RIBO-)
                                                                                                                                                                                                                                             (CHIR )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SANC/) SANCHEZ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  checkpoint protein Chk1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            σ
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                                                                                                                                                                                                                           CHIRON CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGACAGGCCAAAGTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGCCAGGCCAAAGTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 VEGF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                        McSwiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sanchez
                                                                                                                                                                                                                                                                             96US-0584040.
95US-0005974.
                                                                                                                                                                                                                                                                                                                                  96WO-US17480
                                                                                                                                                                                                                            PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             receptor hammerhead ribozyme substrate #438
                                                  218pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             93.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               67.0%;
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                                                                                                                                                                                        Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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Pred. No. 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                        Stinchcomb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ۲,
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present invention describes nucleic acid molecules which modulate synthesis, expression and/or stability of a mRNA encoding 1 or mo

more

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RESULT 21
AAX73045/c
ID AAX730
XX AAX730
XX AAX730
XX Mouse
XX Vascul
KW Vascul
KW Fik-1;
KW foetal
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                                                        The present invention describes nucleic acid molecules which modulate the synthesis, expression and/or stability of a mRNA encoding 1 or more receptors of vascular endothelial growth factor (URGF). A patient (preferably human) having a condition associated with the level of the fms-like tyrosine kinase 1 (flt-1), kinase insert domain containing receptor (KDR) and/or foetal liver kinase 1 (flt-1) (e.g. tumour angiogenesis, ocular diseases, psoriasis and rheumatoid arthritis) can be treated by administering the nucleic acid molecule or the expression vector to the patient. ANX67275 to ANX75725 represent specific examples of nucleic acid molecules from the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acid molecule modulating VEGF receptor(s) mRNA stability - useful for treating e.g. tumour psoriasis, rheumatoid arthritis, etc., in a human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Escobedo J,
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(RIBO-)
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26-OCT-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tumour angiogenesis; psoriasis; rheumatoid arthritis; ocular disease; fms-like tyrosine kinase 1; kinase insert domain containing receptor;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Vascular endothelial growth factor receptor; flk-1; KDR; hammerhead ribozyme; hairpin ribo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-JUL-1999
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95US-0005974.
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                                                                                                                                                                                                       The present invention relates to novel single nucleotide polymorphisms (SNPs) in the human Ran GTPase activating protein 1 (RANGAP1) gene located on chromosome 22q13.2-q13.11, and methods for haplotyping and/or genotyping the RANGAP1 gene. The methods of the invention make use of allele-specific oligonucleotides (ASOs) as probes and primers and/or primer-extension oligonucleotides for detecting the RANGAP1 gene polymorphisms. The polynucleotides and screened compounds are useful for treatment of diseases associated with RANGAP1 activity, such as cancer and other disorders associated with an irregular cell cycle. AAS19743-AAS19820 represent ASO primers for detecting human RANGAP1
                                                                                                                                                     Sequence 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Genotyping human Ran GTPase activating protein 1 gene of individual determining haplotype of individual, involves determining identity of nucleotide pair at specific polymorphic sites for two copies of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; single nucleotide polymorphism; SNP; RANGAP1; haplotyping chromosome 22q13.2-q13.31; Ran GTPase activating protein 1; geno cancer; irregular cell cycle associated disorder; ASO; primer; s
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                                                                                                                                                                                           polymorphisms
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                                                                                                                                                   BP; 3 A; 6 C; 5
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 pulmonary vasocometriction; acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary distress syndrome; pain; cystic fibrosis; pulmonary hypertension; pulmonary vasoconstriction; emphysema; pulmonary hypertension; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer; breast cancer; lung cancer; pancreatic metastasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           which involves the topical administration of an essentially adenosine free antisense oligonucleotide (ON) to the airway epithelium of the subject. The present sequence is an antisense oligonucleotide specific for chymase. The method can be used to treat airway diseases such as cystic fibrosis, asthma, chronic obstructive pulmonary disease, bronchitis and other airway diseases characterised by an inflammatory response. By eliminating adenosine from the antisense ON, its liberation upon antisense degradation is prevented, thereby preventing adenosine-induced bronchoconstriction in patients with hyper-reactive airways.
                                                                                          Antisense oligonucleotide; multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vasoconstriction; inflammation; allergic rhinitis;
                                                                                                                                                                                                                                                          AAX54253 standard; DNA; 14 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 14 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A method for treating airway disease in a subject has been produced, which involves the topical administration of an essentially adenosine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      airway epithelium; adenosine free; cystic fibrosis; obstructive pulmonary disease; bronchitis; ss.
                                                                                                                                                            antisense oligonucleotide fragment
                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                      GGGCCAGGCCAAAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of airway diseases such as asthma - by topically applying -free antisense oligo:nucleotide to airway epithelium of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page
                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nyce JW;
                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                   54.0%;
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RESULT 25
AAF19819/c
ID AAF19819;
XX AAF19819;
AC AAF19819;
XX I-MAR-200
XX I-MAR-200
XX Low adenos
KW Low adenos
KW human; atox
KW human; atox
KW surfactant

14-MAR-2001

(first

entry)

standard;

DNA;

14

ВP

Human chymase polynucleotide fragment #1386.

Low adenosine antisense oligonucleotide, phosphorothicate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic;

respiratory obstruction; pulmonary obstruction; impeded respiration;

pulmonary vasoconstriction; asthma;

surfactant hypoproduction;

밁 Ś

14

GGGCCAGGGCAAGG 1

5 GGGCCAGGCCAAAG

18

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Matches
                Query Match
Best Local
                                                                                                 oligonuclectides (specifically AAX55180-271) can be used for the antisense treatment of diseases and conditions. Typical diseases and conditions are those associated with impaired respiration and inflammation, including lung diseases, pulmonary vasoconstriction, inflammation, allergic rhinitis, acute asthma, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, pulmonary vasoconstriction, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukemias, lymphomas, carcinomas e.g. colon cancer, breast cancer, lung cancer, pancreatic cancer, hepatocellular carcinoma, kidney cancer, melanoma, hepatic mecastases, as well as all types of cancers which may metastasize or have metastasized to the lungs, including breast and prostate cancer.
                                                                                                                                                                                                                                                                                                                The specification describes antisense oligonucleotides (AAX52869-X55271) directed against at least 2 mRNAs selected from target genes, coding and non-coding regions of RNAs corresponding to target genes, gene initiation codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'-end and the juxta-section between coding and non-coding regions and all segments of RNAs encoding proteins associated with one or more diseases, conditions or mixtures. The antisense oligonucleotides may be derived from sequences AAX55272-74. These multiple target
                                                                   Sequence 14 BP; 0 A; 8 C; 3 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 60; 120pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense oligonucleotides used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1999-229400/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-JUN-1998;
17-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    vasoconstriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYEC-) UNIV
 12;
                  Similarity
 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EAST CAROLINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98US-0093972
97US-0059160
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             88
                54.0%;
85.7%;
 0
                  Score
Pred.
 Mismatches
                  10.8;
No. 5
                     59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    treatment of,
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 2
                                Length
 Indels
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 0
Gaps
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CC (I) can have respiratory, bronchodilator, antinfilammatory, analysis, communosuppressive, antiasthmatic, hypotensive and cytostatic activities. CC immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. CC immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. CC immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. CC immunoglobulins and antibodies, antibody receptors, cytokines and contending proteins, adhesion molecules and thair receptors, cytokines and communoglobulins and antibodies, antibody receptors, cytokines and communoglobulins and antibodies, antibody receptors, cytokine and communoglobulins and peripheral nervous and non-nervous system peptide receptors, communities, defensins, growth factors, vasoactive peptides and communities, and peripheral nervous and non-nervous system peptide receptors, binding proteins and malignancy associated proteins. The antisense oligonucleotion which are associated with a disease or condition sepiratory vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome (CRDS), pain, cystic fibrosis (CP), allergies, hinitis (AR), pulmonary cancer. Apfled4 to Apf21543 represent human polynucleotide community of the present invention.
  AXU
                                                             RESULT 26
AAA33697/c
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                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 12
AAA33697
                                       AAA33697 standard; DNA; 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes low adenosine (A) content antisense oligonucleotides and compositions (I) comprising them. In the antisense oligonucleotides the A is replaced by a 'Universal' or alternative base (I) can have respiratory, bronchodilator, antiinflammatory, analysis,
                                                                                                                                                                                                                                                                                                                     Sequence 14 BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Low adenosine (A) content antisense oligonucleotides which do not trigger adenosine receptors during metabolism, useful e.g. for treating cancers and respiratory obstructions -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pulmonary hypertension; emphysema; pulmonary transplantation rejection;
chronic obstructive pulmonary disease; pulmonary infection; bronchitis;
cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 14; Page 250; 1592pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-679539/66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Мусе JW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200062736-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                   14
                                                                                                                                                                                                                                    12;
                                                                                                                                                                                          5 GGGCCAGGCCAAAG 18
                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                GGGCCAGGGCAAGG 1
                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                               invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-0127958
                                                                                                                                                                                                                                                                                                                     A; 8 C; 3
                                                                                                                                                                                                                                                     54.0%;
85.7%;
                                                                                                                                                                                                                                                                                                                     G; 3 T; 0 other;
                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                          Pred.
                                                                                                                                                                                                                                                                           Score 10.8;
                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                          No.
                                                                                                                                                                                                                                                          59;
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                                                                                                                                                                                                                                                                           1.
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                                                                                                                                                                                                                                                                         Length 14;
                                                                                                                                                                                                                                  Indels
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                                                                                                                                                                                                                                  Gaps
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Human; adenosine receptor; low adenosine antisense oligonucleotide; phosphorothioate; impaired respiration; inflammation; allerg; phosphorothioate; impaired respiration; infilammatior; antiinflammatory; allergic disease, bronchoconstriction; inhibitor; antiflammatory; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; lumonary vasoconstriction; asthma; pulmonary vasoconstriction; asthma;
                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-205971/18
                                                                                                                                                                                                                                                                                                                                                                                        Nусе JW
                                                                                                                                                                                                                                                                                                                                                                                                                                 03-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 respiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; COPD;
                                                                                                                                                                                                                                                                                                                                                                                                            (UYEC-) UNIV EAST CAROLINA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200009525-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer; leukaemia; lymphoma; carcinoma; metastasis; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 adenosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                 98US-0095212.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     99WO-US17712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO:1386
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New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstriction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or Claim 18; Page 438; 1343pp; English

aschma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukaemias, lymphomas, carcinomas, and cancers which may metastasise to the lungs, including breast and prostate cancer. The reduction of the adenosine content of the ONs reduces side effects. The A-containing ONs break down with the release of deoxyadenosine which activates adenosine receptors causing bronchoconstriction and inflammation. AAA32311 to AAA35312 represent the nucleotide sequences given in the sequence listing from the present invention, which correspond to SEQ ID NO:1 to 2815, and then the last of the present in the present invention. N.B. Sequences given in the disclosure of the present invention do not match up with their corresponding SEQ ID NO: sequences inflammation. The ON can have antiinflammatory, antiallergic, antiasthmatic, cytostatic and analgesic activities. The compositions are useful for the treatment of diseases associated with inflammation, impaired airways, including lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating e.g. ischaemic conditions, pulmonary vasoconstriction, allergies, The present invention describes a new composition comprising an antisense oligonucleotide (ON) with low adenosine (up to 15%), which targets nucleic acids involved in bronchoconstriction, allergies, and/or in the sequence listing.

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Sequence 14
BP; 0
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C; 3 G; 3 T; 0 other;
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                            Query Match
Best Local S
Matches 12
14
              G
                                    Similarity
GGGCCAGGGCAAGG
             GGGCCAGGCCAAAG 18
                             Conservative
                                   54.0%;
85.7%;
                            0
                                 Score 10.8; 1
Pred. No. 59;
                                          DB
                                          1.
                            2
                                          Length 14;
                             0
                            Gaps
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RESULT 28
AAZ41739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 27
AAZ41523
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                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 11
                                                                                                                                                                                                                                                                                                                                                                                A method has been developed for the amplification of a DNA fragment in which amplification is carried out on the DNA fragments of a number of different DNAs. The method comprises a PCR reaction repeatedly carrying out a heat-denaturing step, a primer annealing step and a polymerase extending step, to amplify the DNA fragments of a plural of different DNAs. The method can detect the existence of a microbe in organic waste. AAZ4124 to AAZ41639 represent PCR primers used in random amplified DOlymorphic DNA arbitrarily primed PCR, for the detection of microbes in organic waste.
                                             Amplification; polymerase chain reaction; PCR; microorganism; compost; detection; pollucant; soil; food; agricultural chemical; polymer; organochlorine; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Microbe; detection; organic waste; arbitrarily pr random amplified polymorphic DNA; amplification;
DE19914461-A1
                       Synthetic
                                                                                             Organic material detecting primer 100
                                                                                                                        20-MAR-2003
21-JAN-2000
                                                                                                                                                             AAZ41739;
                                                                                                                                                                                                                                                                                                                                                 Sequence 12 BP; 3 A; 4 C; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Amplification of a DNA existence of a microbe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31-MAR-1998;
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                                                                                                                                                                                  AAZ41739 standard; DNA; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-626940/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (SAOL )
(NORI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JP11276176-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Microbe detection in organic waste arbitrarily primed PCR primer #100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-JAN-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAZ41523;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Page 9; 40pp; Japanese
                                                                                                                                                                                                                                                                                                 11;
                                                                                                                                                                                                                                                                         9
                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                           waste.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SANYO ELECTRIC CO LTD. ZH NORIN SUISAN SENTAN
                                                                                                                                                                                                                                                 CAGGCCGAAGTC 12
                                                                                                                                                                                                                                                                         CAGGCCAAAGTC 20
                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                      (updated)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               organic waste; arbitrarily primer PCR;
                                                                                                                                                                                                                                                                                                            52.0%;
91.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   fragment
                                                                                                                                                                                                                                                                                                                                                 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                 Pred. No. 50;
); Mismatches
                                                                                                                                                                                                                                                                                                             Score 10.4;
Pred. No. 50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       order to
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RESULT 29
AAC97874
ID AAC97877
XX AAC97877
XX AAC97
XX Prime
XX Prime
XX Prime
XX Prime
XX Prime
XX Prime
XX IP200
XX IP200
XX IP-MA
XX IP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes a novel method for the amplification of DNA comprising (i) preparing many primers with different probabilities of amplification and (ii) simultaneous polymerase chain reaction (PCR) of many different DNA using these primers. The method is used (i) to differentiate between different microorganisms in a mixed population and (ii) to determine presence/absence of an impurity (pollutant) or its concentration, in e.g. soil, foods, compost etc., typically metals, agricultural chemicals, polymers, organochlorine compounds etc. A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   particular use is monitoring composting of organic material. Amplification with many primers produces a lot of information, so reliability of the test is improved, and many samples may be tested
                      WPI; 2001-011047/02
                                                                                                                                                           19-MAR-1999;
                                                                                                                                                                                                                                                                     03-OCT-2000
                                                                                                                                                                                                                                                                                                                       JP2000270867-A
                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                              Primer; amplification; selective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Primer used to illustrate DNA amplification method SEQ ID 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAC97874;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAC97874 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 12 BP; 3 A; 4 C; 4 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                quickly. AAZ41640-Z41855 represent the primers of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel polymerase chain reaction method, microorganisms and for detecting contami
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1999-592157/51
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16-MAR-1999;
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                                                                                                                                                                                                                19-MAR-1999;
                                                                             (SAOL ) SANYO ELECTRIC CO LTD.
(NORI-) ZH NORIN SUISAN SENTAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SAOL ) SANYO ELECTRIC CO LTD. (NORQ ) SOC TECHNO-INNOVATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                           99JP-0076844.
                                                                                                                                                                                                                99JP-0076844
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99JP-0069694.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
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91.7%;
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Pred. No. 50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AGRIC FORESTY & FI
                                                                                GIJUTSU SANGYO
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AAZ81584/c
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Matches
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19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention relates to a method for amplifying a DNA fragment. The method comprises successive repetitions of heat-denaturing, annealing of a primer and an extending step using a DNA polymerase. The method makes use of a cDNA pool in which the primer is one primer or a pair of primer sets and has an amplification probability which allows it to amplify a DNA fragment from a limited number of the cDNAs among the DNA pool (where the limited number is in the range of 1 to 25). Also included in the invention are apparatus used for carrying out the method, a primer and a DNA polymerase and a kit used for amplifying a DNA fragment. The method can be used to amplify a limited number of cDNAs from a pool in which a wide variety of cDNAs are present. Oligonucleotides AAC97775 - AAC97990 represent primers used in an example illustrating the method of
AAZ80767 to AAZ83941 represent tags transcripts that are preferentially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer;
                                                                                                                          Claim 1; Page 80; 219pp;
                                                                                                                                                                                                            prevention
                                                                                                                                                                                                                                                Isolated polynucleotides differentially and non-metastatic breast cancer cells,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-106079/09
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Roberts
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ROBERTS E
SHANKARA
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98US-0089997.
98US-0090039.
98US-0090040.
98US-0090041.
                                                                                                                                                                                                            treatment
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BL.
AS.
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91.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                S
                                                                                                                                                                                                        of cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cell upregulated transcript
                                                                                                                              English.
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Pred. No. 50;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
        corresponding to distinct transcribed in the metastatic breast
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               its apparatus
                                                                                                                                                                                                                                        expressed between metastatic useful for diagnosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1;
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RESULT 31
AAZ86676/c
ID AAZ8666
XX AAZ8666
XX Metast
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XX Human,
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CC AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts

CC that are preferentially transcribed in the primary or non-metastatic

CC breast tumour tissue (i.e. are downregulated in metastatic breast tumour

CC cells). These transcripts can be used for diagnosis, prognosis,

CC monitoring and treatment of breast cancer, particularly where metastatic.

CC Diagnosis is by standard immunoassays or hybridisation/amplification

CC reactions. Compounds that modulate expression of the transcripts are

CC promoters from the transcripts are used to direct expression, in selected

CC cell types, of e.g. therapeutic genes (also ribozymes or antisense

CC sequences), particularly an antigen-encoding sequence for use in gene or

CC cell-based vaccines. Polypeptides encoded by the transcripts are also

CC useful in vaccines; for diagnosing breast cancer and for raising

CC specific antibodies (Ab). Ab are used to detect the polypeptides or as

CC therapeutic agents. Host cells that produce the polypeptides can be used

CC constituted and isolate populations of educated, antigen-specific immune
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Best Local S
Matches 10
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                                                               prevention and
                                                                                     Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis,
                                                                                                                                                                                              Roberts
                                                                                                                                                                                                                                                             (GENZ )
(ROBE/)
                                                                                                                                                                                                                                                                                                                                                                                    19-JUN-1998;
19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Metastatic breast tumour cell downregulated transcript tag
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                                                                                                                                                                                                                                                                                                                           19-JUN-1998
                                                                                                                                                                                                                                                                                                                                            19-JUN-1998;
19-JUN-1998;
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                                                                                                                                                    2000-106079/09.
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                                                                                                                                                                                                                                   GENZYME CORP.
ROBERTS B L.
SHANKARA S.
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                                                                                                                                                                                            BL,
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                                                                                                                                                                                            Shankara
                                                                 treatment of cancer
                                                                                                                                                                                                                                                                                                                                            98US-0090039
98US-0090040
                                                                                                                                                                                                                                                                                                                                                                                    98US-0089853
98US-0089997
                                                                                                                                                                                                                                                                                                                                                                                                                                                   99WO-US13647.
                                                                                                                                                                                                                                                                                                                           98US-0090041
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     vaccine; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11
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Pred. No.
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38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       breast cancer; tag; pr
ne therapy; anticancer;
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Claim 1;

Page

214;

219pp; English

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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected call types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or call-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines, for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for
The present invention describes a method of identifying the type of cell in a sample, involving determining which of the sequences AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1570/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis,
                                                                                                                                                      New isolated polynucleotides, useful type, such as cancer cell, comprises
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 10 BP;
                                                                                                        Claim
                                                                                                                                                                                                                                          Velculescu
                                                                                                                                                                                                                                                                                                          24-NOV-1999;
                                                                                                                                                                                                                                                                                                                                        21-NOV-2000; 2000WO-US31922
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human ubiquitously expressed transcriptome sequence SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAH64570;
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    adoptive immunotherapy.
                                                                                                                                                                                                                                                                         (UYJO ) UNIV JOHNS HOPKINS.
                                                                                                   13; Page 71; 94pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         transcriptome;
diagnosis; cell
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                                                                                                                                      cell types
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                                                                                                                                                                                                                                        Vogelstein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ome; gene expression pattern; cancer; cell specific gene expression; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entry)
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                                                                                                                                                        comprises
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Т;
                                                                                                                                                                                                                                          Kinzler
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                                                                                                                                                    for identifying specific transcriptomes expressed
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RESULT 34
ABZ72939
ID ABZ72

ABZ72939

standard; RNA;

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RESULT 33
AAH64571/C
ID AAH645
XX Human
XX Human
XX Human
XX Homo s
XX Cancer
XX Cancer
XX 21-NOV
XX 21-NOV
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Matches 10
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                                                                                                                                          The present invention describes a method of identifying the type of cell in a sample, involving determining which of the sequences AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce cancer specific gene expression, standardise expression and restore the function of a diseased cell or tissue. The present sequence is one of the transcriptomes described in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cancer specific gene expression, standardise expression and restore the function of a diseased cell or tissue. The present sequence is one of the transcriptomes described in the exemplification of the invention.
                                                                                                                   Sequence 10
                                                                                                                                                                                                                                                                                                            particular cell types -
                                                                                                                                                                                                                                                                                                                          New isolated polynucleotides, useful for identifying specific type, such as cancer cell, comprises transcriptomes expressed
                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-367706/38.
                                                                                                                                                                                                                                                                                                                                                                                                     Velculescu
                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; transcriptome; gene expression pattern; cancer; drug screening; cancer diagnosis; cell specific gene expression; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human ubiquitously expressed transcriptome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAH64571;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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                              2 TCCGGGCCAG
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                                                                        Similarity
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TCCGGGCCAG
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                                                                                                                 BP; 1 A; 4 C; 4 G; 1 T; 0 other;
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                                                           Conservative
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                                                                                                                                                                                                                                                                               71; 94pp;
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                                                                                                                                                                                                                                                                                                                                                                                                  Vogelstein B,
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                                                                      100.0%;
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                                                                      Score 10;
Pred. No.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                     Kinzler KW
                                                           Mismatches
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                                                                      DB 1;
38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence SEQ ID NO: 1411.
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in
                                                        Gaps
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S
                                                                                                                                                                              (AAV) rectored ribozyme composition (1). (1) comprises: (a) at least a first ribozyme that specifically cleaves an mRNA encoding a protein, polypeptide, or peptide selected from the group of rod opsin, iNOS, RDS/peripherin, VEGFR1, VEGFR2, adenosine A-2B receptor, IGF-1, integrin alpha 1, integrin alpha 5, or integrin alpha V; (b) a vector comprising a polynucleotide encoding the ribozyme, where the polynucleotide operably positioned downstream of at least a first polynucleotide operably positioned downstream of at least a first comprising the ribozyme or the polynucleotide; (c) a viral particle comprising the ribozyme or the polynucleotide; or (e) a host cell comprising the ribozyme or the polynucleotide; or (e) a host cell for decreasing the amount of mRNA encoding a selected polypeptide in a retinal cell of a mammalian eye, comprising providing to the eye the composition described above, and for a time effective to specifically cleaves the mRNA in the cell. (I) has ophthalmological activity, and can be used in gene therapy. (I) can be used for treating a disease or retinal dis
                                                                           Matches
                                                                                                     Query Match
Best Local :
                                                                                                                                                                                                                                               dysfunction, (diabetic) retinopathy, or (age-related) macular degeneration. (I) is also useful for manufacturing a medicament for treating the diseases mentioned above, including autosomal dominant retinitis or a blood-retinal barrier dysfunction. (I) can also be useful for treating, decreasing the severity, or ameliorating the symptoms of a pathological condition, e.g. atrophic or pigmented lesions of the eye, blindness, a reduction in central or peripheral vision, or a reduction in total vision. ABZ7263 to ABZ72953 represent sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A recombinant adeno-associated virus-vectored ribozyme composition, useful for treating a disease or dysfunction of the mammalian eye eretinal disease, e.g. diabetic retinopathy or age-related macular
                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                  exemplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 5; Page 73; 115pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-111880/10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-MAY-2002; 2002WO-US13679.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hairpin ribozyme; hammerhead ribozyme; ribozyme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rod opsin hammerhead ribozyme oligonucleotide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              present invention describes a recombinant adeno-associated virus
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6 GGCCAGGCCA 15
                                                                                                     Similarity
                                                                                                                                                                              12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Shaw LC,
                                                                              Conservative
                                                                                                                                                                              BP; 3
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                                                                                                                                                                      A; 4 C; 4 G; 1 U; 0 other;
                                                                                                                                                                                                                             the present
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Pred, No. 61;
                                                                                                                                                                                                                               invention.
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AAA04950/ ID AAAC XX

AAA04950 standard;

DNA;

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RESULT 36

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AAV63772/c
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XX
AC AAV637
XX
AC AAV637
XX
DT 26-APR
XX
Human
XX
Hy JAGGED
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Matches
                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                      This nucleotide corresponds to a portion of exon 23 of the human JAGGED1 gene that is mutated in affected individuals of an Alagille Syndrome (AS) family. The mutation involves insertion of 'GTGGC' residues at a position corresponding to nucleotide 3102 (see AAV63773) of an hJAGGED1 cDNA (see AAV63753). The mutation causes product (see AAW67904). It was detected by heteroduplex mobility analysis (see AAW6376-68 and AAV84310-19). hJAGGED1 (see also AAW87994) is a novel Notch ligand capable of inhibiting the differentiation of haematopoietic progenitor cells. Methods are provided for the diagnosis of AS based on detection of disease-associated mutations or truncated hJAGGED1 gene products.
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progenitor cell; haematopoiesis; cell differentiation; mutant;
Alagille syndrome; leukaemia; lymphoma; diagnosis; therapy; ss
                                                                                                                                                                                                                                       Sequence 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 5; Fig 8; 101pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        state, e.g. for haematopoietic reconstitution
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New Jagged peptides for inhibiting differentiation of progenitor cells - also used for maintaining these cells in undifferentiated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-081220/07
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26-APR-1999
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                                                                                                                                                  Local
                                                        5 GGGCCAGGCCAAA 17
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                                                                                                                                                  Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GGCCAGGCCA
                                                                                                                                                                                                                                       BP; 0 A; 4 C; 6 G; 3 T; 0 other;
                                                                                                                   Conservative
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/note= "5-base insertion"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                               49.0%;
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                                                                                                                                                                               Score 9.8;
                                                                                                                                                  Pred. No.
                                                                                                                      Mismatches
                                                                                                                                                  81;
                                                                                                                                                                               DB 1;
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                                                                                                                                                                           Length 13;
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RESULT 37
AAA05213/c
ID AAA052
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XXXXXXX
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                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 11; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; Tenascin-C; extracellular matrix protein; phosphorothioate; antisense oligonucleotide; inhibition; exon deletion; therapy; cellular development; differentiation; translation; ss.
                                               18-MAY-2000
                                                                                                                AAA05213 standard; DNA; 13
                                                                                                                                                                                                                                                                                                                                     Sequence 13 BP; 0 A; 5 C; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Preparing antisense oligodeoxynucleotides (ODNs) and long antisense sequences useful for blocking translation of a specific isoform of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fillmore H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
          Tenascin-C phosphorothioate antisense oligonucleotide SEQ ID NO:502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23; Page 102; 177pp; English
                                                                                                                                                                                                    13
                                                                                                                                                                                                                                                                                                                                                                                    sequence listing but not mentioned further within the
                                                                                                                                                                                                                                  δ
                                                                                                                                                                                                    GGCCAAGCCGAAG 1
                                                                                                                                                                                                                                  GGCCAGGCCAAAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein
                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Broaddus WC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      9808-0094255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99WO-US16632
                                                                                                                                                                                                                                                                                  49.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gillies GT,
                                                                                                                                                                                                                                                                                                                                     G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                    <u>..</u>
                                                                                                                                                                                                                                                                                  Score 9.8;
Pred. No. 8
                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                   81;
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                                                                                                                                                                                                                                                                  2
                                                                                                                                                                                                                                                                                                  Length 13;
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ABC48592/c
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AT 21-FEB
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OS SNP; s
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes a method for preparing an antisense oligodeoxynucleotide (ODN) sequence for blocking translation of a specific protein isoform that can be expressed as a number of different isoforms. AAAO4712 to AAAO5243 represent specifically claimed phosphorothioate antisense ODNs for blocking translation of Tenascin-C using the method of the invention. The method is useful for preparing an ODN sequence for blocking translation of a specific isoform of Tenascin-C protein. The method is also useful for blocking translation of a specific family of isoforms of a protein. The method can also be performed by producing a long antisense expression vector encoding a long antisense expression vector encoding a long antisense expression vector section of designing models for studying cellular development and differentiation. The method permits selective inhibition of the translation of protein isoforms, which occur as a result of alternative splicing. AAAO5244 represent an oligonucleotide from the present invention, which is given
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; Tenascin-C; extracellular matrix protein; phosphorothioate; antisense oligonucleotide; inhibition; exon deletion; therapy; cellular development; differentiation; translation; ss.
                                    Homo sapiens
                                                                                                                                                                                                                          Oligonucleotide SEQ ID NO 48609 for detecting SNP TSC0013829
                                                                                                                                                                                                                                                                                         21-FEB-2002
                                                                                                                                                                                                                                                                                                                                                        ABC48592
                                                                                                                                                                                                                                                                                                                                                                                                                       ABC48592 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Preparing antisense oligodeoxynucleotides (ODNs) and long antisense RNA sequences useful for blocking translation of a specific isoform of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fillmore H,
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence listing but not mentioned further within the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6 GGCCAGGCCAAAG 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GGCCAAGCCGAAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 0 A; 5 C; 4 G; 4 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                         (first
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RESULT 39
ABC48593
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation.

ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH02073 represent the oligomers described in the invention.

NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fip.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Set of oligonucleotides, useful for diagnosis and cell typing, designed to detect single nucleotide polymorphisms and cytosine methylation status -
                                                                                                                                                                                                                                                                                                                                                                                                                                               peptide nucleic
central nervous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; peptide nucleic acid; cytosine methylation; cardiovascular;
WPI; 2001-657177/75
                                                    Olek A,
                                                                                                                                                                                                                         06-APR-2001; 2001WO-IB00713.
                                                                                                                                                                                                                                                                                   18-OCT-2001.
                                                                                                                                                                                                                                                                                                                                    WO200177384-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-APR-2000; 2000DE-1019173
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                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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                                                                                                              (EPIG-)
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                                                                                                              EPIGENOMICS AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EPIGENOMICS AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                    Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CCAGGCCAAAGTC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Piepenbrock C, . Berlin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                     2000DE-1019173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                               system; gastrointestinal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ID NO 48610 for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             49.0%;
                                                       Berlin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 9.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             No. 81;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP TSC0013829.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer; cns;
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Best Local S
Matches 11
This invention describes a novel non-MRL healer mouse (M) having at least one quantitative trait locus selected from those given in the specification, exhibiting an enhanced healing response to a wound compared to mice (m) without the locus. The invention describes a numethod of identifying a gene involved in enhanced wound healing by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99889, ABH00010-ABH99989 and ABI000100-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the prin specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Wound healing; non-MRL healer mouse; quantitative trait locus; QTL healing response; microsatellite marker; treatment; central nerve; peripheral nerve; nerve injury; SAGE tag; murine; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell designed to detect single nucleotide polymorphisms and methylation status \,
                                                                                              Claim
                                                                                                                                           New mammalian model for enhanced wound healing
                                                                                                                                                                          WPI; 1999-494533/41
                                                                                                                                                                                                                                                                   26-AUG-1998;
                                                                                                                                                                                                                                                                                28-SEP-1998;
13-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                               19-AUG-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Murine MRL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAZ19014;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID 48610; 29pp + Sequence Listing; German.
                                                                                                                         identifying
                                                                                                                                                                                                                                                                                                                                12-FEB-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CCAGGCCAAAGTC 20
                                                                                             Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SAGE tag
                                                                                                                         enhanced wound healing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                98US-0102051.
98US-0074737.
98US-0097937.
                                                                                             74; 136pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 9.8; D
Pred. No. 81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                           genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and cell typing,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2
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RESULT 41
AAX55034/c
ID AAX550
AC AAX550
AC AAX550
AC AAX550
AC AAX550
AC CEBP-
XX Antise
KW Impair
KW Impair
KW pulmon
KW colon
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Best Local S
Matches 10
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non-coding regions of RNAs corresponding to target genes, gene initiation codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'-end and the juxta-section between coding and non-coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  pulmonary hypertension; pulmonary vasoconstriction; emphysema;
chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma;
colon cancer; breast cancer; lung cancer; pancreatic cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense oligonucleotide; multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vasoconstriction; inflammation; allergic rhinitis; acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis;
                                                                                                    The specification describes antisense oligonucleotides directed against at least 2 mRNAs selected from target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-JUN-1998;
17-SEP-1997;
                                                                                                                                                                                                     Disclosure; Page 70; 120pp;
                                                                                                                                                                                                                                                                        vasoconstriction
                                                                                                                                                                                                                                                                                               New antisense oligonucleotides used in treatment of, e.g. pulmonary
                                                                                                                                                                                                                                                                                                                                                                      WPI; 1999-229400/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hepatocellular carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYEC-) UNIV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EAST CAROLINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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97US-0059160.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     47.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    kidney cancer; melanoma; hepatic metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         σ
                                                                                                                                                                                                     English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred.
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                                                                                                           genes,
                                                                                                                                     (AAX52869-X55271)
      and non-coding
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                                                                                                 coding and
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RESULT 42
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inflammation, including lung diseases, pulmonary vasoconstriction, inflammation, allergic rhinitis, acute asthma, allergies, asthma, inpeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, pulmonary vasoconstriction, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukemias, lymphomas, carcinomas e.g. colon cancer, breast cancer, lung cancer, pancreatic cancer, hepatocellular carcinoma, kidney cancer, melanoma, hepatic metastasses, as well as all types of cancers which may metastasize or have metastasized to the lungs, including breast and prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           regions and all segments of RNAs encoding proteins associated with one or more diseases, conditions or mixtures. The antisense oligonucleotides may be derived from sequences MAX55272-74. These multiple target oligonucleotides (specifically AAX552180-271) can be used for the antisense treatment of diseases and conditions. Typical diseases and conditions are those associated with impaired respiration and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Low adenosine antisense oligonucleotide; phosphorothioate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory;
                                            Claim
                                                                                         Low adenosine (A) content antisense oligonucleotides which trigger adenosine receptors during metabolism, useful e.g.
                                                                                                                                         WPI; 2000-679539/66
                                                                                                                                                                         Nyce JW
                                                                                                                                                                                                                                                      06-APR-1999;
                                                                                                                                                                                                                                                                                    24-MAR-2000; 2000WO-US08020
                                                                                                                                                                                                                                                                                                                    26-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                              surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pain; cystic fibrosis; allergic rhinitis; pulmonary hypertension; emphysema; pulmonary transplantation rejection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 11 BP; 0 A; 6 C; 5
                                                                                                                                                                                                                                                                                                                                                   WO200062736-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                   chronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             respiratory obstruction; pulmonary obstruction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human C/EBP polynucleotide fragment #2170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF20603
                                                                                                                                                                                                      (UYEC-) UNIV EAST CAROLINA (NYCE/) NYCE J W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11
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                                                                                                                                                                                                                                                                                                                                                                                                                  88.
                                                                                                                                                                                                                                                                                                                                                                                                                               obstructive
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cceecceec
                                           Page
                                                                          respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first
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                                           1592pp;
                                                                                                                                                                                                                                                                                                                                                                                                                               pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry)
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                                                                             obstructions
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                                           English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0,:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 9.4;
Pred. No. (
                                                                                                                                                                                                                                                                                                                                                                                                                                 disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                               pulmonary infection; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              impeded respiration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                           do not
for treating
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The present

invention describes low adenosine (A) tides and compositions (I) comprising

content them. In

antisense the antis

antisense

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RESULT 43
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                               Determining Tm range for several degenerate primers with a fixed-sequence and a degenerate-sequence portion for use in polymerase chain reaction amplification by identifying a specific sequence in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pulmonary transplantation rejection, pulmonary infections, bronchitis, and/or cancer. AAF18434 to AAF21543 represent human polynucleotide fragments and antisense oligonucleotides used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    condition selected from pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome (RDS), pain, cystic fibrosis (CF), allergic rhinitis (AR), pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD), hypertension, emphysema, chronic obstructive pulmonary disease (COPD).
                                                               nucleic
                                                                                                                                                                                                                                                                                                                                                                                                       06-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-APR-2000; 2000WO-US08962
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer; nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAC63229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAC63229 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 11 BP; 0 A;
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                                                                                                                                                                                                                                                                                                                                          (GENO-) GENOME
                                                                                                                                                                                                                  2000-656235/63
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        invention.
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                                                         template
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                       TECHNOLOGIES LLC
                                                                                                                                                                                                                                                                                                                                                                                                    99US-0127891.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  #2 used in a method for primer selection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               acid amplification; melting temperature; T_m; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               47.0%;
90.9%;
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C;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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Disclosure; Fig 2; 34pp;

The present invention describes a new composition comprising antiense oligonuclectide (ON) with low adenosine (up to 15%) targets nucleic acids involved in bronchoconstriction, right inflammation. The ON can have antiinflammatory, antiallergic, antiasthmatic, cytostatic and analgesic activities. The compouseful for the treatment of diseases associated with inflamma

allergies, and/or compositions are

which

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RESULT 44
AAA34481/c
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a method for selecting PCR primers for nucleic acid amplification. The method comprises determining the melling temperature (T_m) range for degenerate oligonucleotide primers with a fixed-sequence portion (FS) and a degenerate-sequence portion (DS) by searching known portion of a nucleic acid template for a sequence complementary to a desired FS of a primer. Nucleotide base pairs flanking or interspersed between the sequence complementary to a DS of one of the primers are detected and T m is calculated. The method of the present invention allows primers which produce more efficient DNA amplification to be produced. The present sequence is a primer This sequence was used to exemplify the occurence of a primer with a FS of 6 base pairs (CGGCCC) within a template. The remaining 5 base pairs make up the DS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; adenosine receptor; low adenosine antisense oligonucleotide; phosphorothicate; impaired respiration; infilammation; allergy; allergic disease; bronchoconstriction; inhibitor; antinflammatory; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; antiallergic; actiasthmatic; cytostatic; analgesic; impaired airway; lung disease; ischaemic condition; pulmonary vasoconstriction; asthma;
                                                                                                                                                                   New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstriction, inflammation, allergies, asthma, hypertension,
                                                                                                         Disclosure;
                                                                                                                                                      bronchitis,
                                                                                                                                                                                                                       WPI; 2000-205971/18
                                                                                                                                                                                                                                                                                                                     03-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                   03-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                  24-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                  WO200009525-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pulmonary hypertension; chronic cancer; leukaemia; lymphoma; can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             respiratory distress syndrome; pain; cystic fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human adenosine receptor related polynucleotide SEQ ID NO:2170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA34481 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 11 BP; 1 A; 6 C; 4 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                    (UYEC-) UNIV EAST CAROLINA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CGGCCCAGGCC 11
                                                                                                      Page
                                                                                                                                                        emphysema,
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                                                                                                                                                                                                                                                                                                                                                   99WO-US17712
                                                                                                        537;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              47.0%;
90.9%;
                                                                                                      1343pp; English
                                                                                                                                                        respiratory distress
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 9.4; D
Pred. No. 66;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           obstructive pulmonary disease; COPD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                        syndrome,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               88
                                                                                                                                                        ischemia or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             emphysema;
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RESULT 45
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ID ABV623
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XX ABV623
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PN W02002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local :
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asthma,
The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     impaired airways, including lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating
                                                                                                                                                                                                                                                                                                                                                                             In vitro identification of homeostasis and identifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-JAN-2001; 2001DE-1000127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABV62380 standard; cDNA; 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 11
                                                                                                                                                                                                                                                                       Disclosure; Page 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-590638/63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Petersohn D,
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Pred. No. 66;
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                                                                                                                                                                                                                                                                                           Disclosure; Page 147; 1345pp;
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RESULT 47
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YX 11-JU
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                                                                                    AAX55033;
                                                                                                                                   AAX55033 standard; DNA; 12
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                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                             Length 11;
                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             scleroderma;
                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                            0;
AAF20602/c
ID AAF206
XX
AC AAF206
XX
                                                                                                                                   RESULT 49
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AAF20602

AAF20602 standard; DNA; 12

ВP

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В
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                                                                                                                                        Matches
                                                                                                     Query Match
                                                                                                                                                                     inflammation, including lung diseases, pulmonary vasoconstriction, inflammation, allergic rhinitis, acute asthma, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, pulmonary vasoconstriction, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukemias, lymphomas, carcinomas e.g. colon cancer, breast cancer, lung cancer, pancreatic cancer, hepatocellular carcinoma, kidney cancer, melanoma, hepatic metastases, as well as all types of cancers which may metastasize or have metastasized to the lungs, including breast and prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                      non-coding regions of RNAs corresponding to target genes, gene initiation codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'-end and the juxta-section between coding and non-coding regions and all segments of RNAs encoding proteins associated with one or more diseases, conditions or mixtures. The antisense oligonucleotides
                                                                                                                                                                                                                                                                                                                                    may be derived from sequences AAX55272-74. These multiple target oligonucleotides (specifically AAX55180-271) can be used for the antisense treatment of diseases and conditions. Typical diseases and conditions are those associated with impaired respiration and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary hypertension; pulmonary seconstriction; emphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer; hepatic cancer; kidney cancer; melanoma; hepatic metastasis;
                                                                                                                                        Sequence 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The specification describes antisense oligonucleotides (AAX52869-X55271) directed against at least 2 mRNAs selected from target genes, coding and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-229400/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nyce JW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-JUN-1998;
17-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9913886-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 70; 120pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       vasoconstriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         pulmonary vasoconstriction; inflammation; allergic rhinitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C/EBP-beta antisense oligonucleotide fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (UYEC-) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       impaired respiration;
                                                                                      Local
12
                                 w
                                                                    Similarity
10; Conserv
                               cceeeccaeec
                                                                                                                                        BP; 0
                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EAST CAROLINA
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97US-0059160
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                                                                                                                                        Α,
                                                                                  47.0%;
90.9%;
                                                                                                                                        7 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       inflammation;
                                                                                                                                        ഗ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          multiple
                                                                     0
                                                                                                                                        G; 0 U; 0 other;
                                                                                    Score 9.4;
Pred. No. 8
                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     target; antis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     in treatment of, e.g. pulmonary
                                                                                      82
                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antisense treatment;
                                                                     ۳.
                                                                                                   Length 12;
                                                                     Indels
                                                                    0
                                                                    Gaps
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Human C/EBP polynucleotide fragment #2169

(first entry)

enosine antisense oligonucleotide; phosphorothioate; airway disorder; bronchoconstriction; lung inflammat

lung inflammation;

allergy;

surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic; respiratory obstruction; pulmonary obstruction; impeded respiration; surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pain; cystic fibrosis; allergic rhinitis; pulmonary hypertension; emphysema; pulmonary transplantation rejection; chronic obstructive pulmonary disease; pulmonary infection; bronchitis;

Homo sapiens cancer; ss.

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CC immunosuppressive, antiastimatic, hypotensive and cytostatic activities. CC The antisense oligonucleotides and (I) can be used to down-regulate the CC expression and or activity of target polypeptides associated with CC lung/respiratory disorders and malignancies, such as stimulating and activating peptide factors and transmitters, transcription factors, CC immunoglobulins and antibodies, antibody receptors, cytokines and CC elemokines, endogenously produced specific and non-specific enzymes, binding proteins, adhesion molecules and their receptors, cytokine and CC chemokine receptors, adenosine receptors, bradykinin receptors, central concrivous system (CNS) and peripheral nervous and non-nervous system peptide cransmitters, defensins, growth factors, vasoactive peptides and CC receptors, binding proteins and malignancy associated proteins. The CC including respiratory obstruction (especially pulmonary obstruction and/or lung inflammation, allergy(ses) condition selected from pulmonary vasoconstriction, inflammation, condition selected for pulmonary vasoconstriction, inflammation, condition selected for pulmonary vasoconstriction, inflammation, conditions selected for pulmonary vasoconstriction, selected for pulmonary vasoconstric
Query Match
Best Local Similarity
Matches 10; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes low adenosine (A) content antisense oligonucleotides and compositions (I) comprising them. In the antisense oligonucleotides the A is replaced by a 'Universal' or alternative base (I) can have respiratory, bronchodilator, antiinflammatory, analgesic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (UYEC-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 14; Page 264; 1592pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancers and respiratory obstructions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-679539/66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nyce JW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Low adenosine (A) content antisense oligonucleotides which trigger adenosine receptors during metabolism, useful e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-MAR-2000; 2000WO-US08020
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                                                                                 BP;
  Conservative
                                                                                                                                invention.
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                                                                                   ð
                 47.0%;
                                                                                   7 C; 5
0,
                                                                                 G; 0 U; 0 other;
Score 9.4; DB 1;
Pred. No. 82;
0; Mismatches 1
1;
                                       Length 12;
  Indels
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Gaps
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RESULT 50
AAA34480/c
ID AAA344
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KW Ph
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-205971/18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            respiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; COPD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human adenosine receptor related polynucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-JUL-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-FEB-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antiallergic;
lung disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         phosphorothicate; impaired respiration; inflammation; altergy;
allergic disease; bronchoconstriction; inhibitor; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; adenosine receptor; low adenosine antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     leukaemia;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ischaemic condition; pulmonary vasoconstriction; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antiasthmatic; cytostatic; analgesic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98US-0095212.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        lymphoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  impaired airway;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NO:2169
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New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstriction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or

Disclosure; Page 537; 1343pp; English

cc targets nucleic acids involved in bronchoconstriction, allergies, and/or cargets nucleic acids involved in bronchoconstriction, allergies, and/or criminal ammation. The ON can have antiinflammatory, antiallergic, antiasthmatic, cyrostatic and analgesic activities. The compositions are useful for the treatment of diseases associated with inflammation, compaired airways, including lung disease and diseases whose secondary ceffects afflict the lungs of a subject. They can be used for treating e.g. ischaemic conditions, pulmonary vasoconstriction, allergies, casthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, emphysema, chronic obstructive grimonary disease (COPD), and cancers such as leukaemias, lymphomas, carcinomas, and cancers which may metastasise to the lungs, including breast and prostate cancer. The reduction of the adenosine content of the ONs reduces side effects. The A-containing ONs break down with the crelease of deoxyadenosine which activates adenosine receptors causing bronchoconstriction and inflammation. AAA32313 to AAA35312 represent the nucleotide sequences given in the sequence listing from the present invention, which correspond to SEQ ID NO:1 to 2815, and then the last CI appears to AAA33932 to AAA33932 represent the sequences. SEQ ID NO:11 to 1680 (AAA32333 to AAA33992) are specifically claimed ONs from the present convention. N. B. Sequences given in the disclosure of the present convention. N. B. Sequences given in the disclosure of the present convention. N. B. Sequences given in the disclosure of the present convention. N. B. Sequences given in the disclosure of the present convention. N. B. Sequences given in the disclosure of the present convention. N. B. Sequences given in the disclosure of the present convention. N. B. Sequences given in the disclosure of the present convention. N. B. Sequences given in the sequences given in the disclosure of the present convention. N. B. Sequences given in the green convention and convention antisense oligonucleotide (ON) with present invention describes a new composition comprising a isense oligonucleotide (ON) with low adenosine (up to 15%). an which

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ARESULT 51
AAZ79191/C
ID AAZ791
AC AAZ71
AC AAZ791
AC AAZ791
AC AAZ791
AC AAZ71
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Matches 10
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19-JUN-1998;
19-JUN-1998;
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                       WPI; 2000-106077/09
                                                             Roberts BL,
                                                                                                                                                                                                                                                                                                         19-JUN-1998;
19-JUN-1998;
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19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytotoxic T-lymphocyte; tumour antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SAGE tag; serial analysis of gene expression; antigen-presenting cell; APC; monocyte-derived dendritic cell; differential gene expression; immunostimulatory cofactor; costimulatory factor; CTL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human dendritic cell SAGE tag,
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                                                                                                                                                                                                                                                                                        19-JUN
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                                                                                                    GENZYME CORP.
ROBERTS B L.
SHANKARA S.
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                                                             Shankara
                                                                                                                                                                                                                                            98US-009004

98US-009007

98US-009007

98US-009007

98US-009007

98US-009007

98US-009007
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98US-0090080.
98US-0111715.
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98US-0089999
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98US-0089853
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90.9%;
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Isolated polynucleotides differentially expressed in antigen-presenting cells, useful in gene vaccines against cancer -

Page 111; 130pp; English

context transcripts correspond to novel genes in department of the continuitation of the cytotoxic immune response, particularly against tumour cells. Tumour antigen presentation via the MHC (major histocompatibility complex) and subsequent recognition by T-cell receptors is alone complex) and subsequent recognition by T-cell receptors is alone complex) and subsequent recognition by T-cell receptors is alone complex) and subsequent recognition by T-cell receptors is alone complex) and subsequent recognition by T-cell receptors is alone complex to activate a robust cytotoxic immune response that can complex that the tumour cells, immunostimulatory cofactors also being required complex that the complex that can be used in vaccines to induce an immune response, particularly against a tumour antigen; to modulate the genotype of an APC; to screen complex that modulate expression of differentially expressed genes in an application primers for the composis, prognosis and monitoring of diseases related to abnormal compression of these genes. Detection of the dendritic cell containing them are used in active immunotherapy (or to stimulate correction of a population of antigen specific effector cells) and containing them are used in gene therapy. Co-administration of containing them are used in gene therapy. Co-administration of containing them are used in gene therapy. Co-administration of containing them are used in gene therapy. Co-administration of containing them are used in gene therapy. Co-administration of containing them are used in gene therapy. Co-administration of containing them are used in gene therapy. Co-administration of containing them are used in gene therapy. Co-administration of containing them are used in gene therapy. Co-administration of containing them are used in gene therapy. Co-administration of containing them are used in gene therapy. Co-administration of containing the containing them are used in gene therapy. Co-administration of containing the containing the containing the containing the immunostimulatory cofactor proteins which are preferentially or differentially expressed in monocyte-derived dendritic cells compared with monocytes. Some of the transcripts correspond to known genes or ESTs (expressed sequence tags) which were previously unknown to be preferentially or differentially expressed in dendritic cells, while Sequences AAZ77573-Z79709 represent SAGE (serial analysis of gene expression) tags used to identify mRNA transcripts encoding recruitment of immune effector cells.

Sequence 10 BP; 0 A; 3 C; 2 G; 5 T; 0 other;

Query Match Best Local S Matches 9 10 AGGCCAAAG 18 l Similarity 9; Conserv Conservative 45.0%; 100.0%; 0 Score 9; Pred. No. Mismatches DB 1 0 Indels 0 Gaps 0

Length 10;

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В 10 AGGCCAAAG 2

ВP

Metastatic breast tumour cell upregulated transcript tag #1562.

RESULT 52
AAZ82338/c
ID AAZ82328 standard; DNA; 10 BP
XX
AC AAZ82328;
XX
DT 07-APR-2000 (first entry)
XX
DE Metastatic breast tumour cell
XX
DE mon-metastatic breast tumour
XW
Antimetastatic; vaccine; diag
XX
PN W09965928-A2.
XX
PD 23-DEC-1999.
XX
PD 23-DEC-1999.
XX
PF 18-JUN-1998; 98US-0089853.
PR 19-JUN-1998; 98US-0089997. antimetastatic; vaccine; diagnosis; ss. non-metastatic breast tumour tissue; Human; metastatic breast tumour tissue; breast cancer; tag; primer; gene therapy; anticancer;

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RESULT 53
AAZ85483
ID AAZ85
XX AZ85
AC AAZ85
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Best Local S
Matches 9
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19-JUN-1998;
19-JUN-1998;
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AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis,
                                                                                                                                                                                                                                                                Human; metastatic breast tumour tissue; breast cancer; non-metastatic breast tumour tissue; gene therapy; antimetastatic; vaccine; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-APR-2000
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                                                                                          23-DEC-1999.
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                               18-JUN-1999;
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98US-0090040.
98US-0090041.
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anticancer;
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XX AAZ864
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                                                                                                                                      Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer;
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SHANKARA
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CC AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, compitoring and treatment of breast cancer, particularly where metastatic Diagnosis is by standard immunoassays or hybridisation/amplification creations. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising compended to a particularly and the sequence of the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune cells, e.g. cyvotoxic T lymphocytes, and these used for advantive immunerharany.
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                                                                       Human; metastatic breast tumour tissue; non-metastatic breast tumour tissue; ger antimetastatic; vaccine; diagnosis; ss.
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                                                                                                                                  Metastatic breast tumour cell downregulated transcript
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                                            Homo sapiens
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ROBERTS B L.
SHANKARA S.
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98US-0090039.
98US-0090040.
98US-0090041.
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                                                                                                                                                                                                                                                                                                                                                              red. No. 64
Mismatches
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                                                                        ; gene therapy; anticancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                         other;
                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                   breast cancer; tag; primer;
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RESULT 56
ABL52193/c
ID ABL52193 standard; DNA; 10
XX
AC ABL52193;
XX
DT 12-JUL-2002 (first entry)
XX
DT 12-JUL-2002 (forst entry)
XX
Cos Human PER1 preferred oligo
XX
Cos Human; period (Drosophila)
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Cos Human; period (Drosophila)
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Cos Homo sapiens.
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XX
Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour rells).

CC AAAB3942 to AAZB6677 represent tags corresponding to distinct transcripts (that are preferentially transcribed in the primary or non-metastatic breast tumour cells).

CC that are preferentially transcribed in the primary or non-metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, to monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are promoters from the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense captences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising constitutions of educated, antigen-specific immune effecter cells, e.g. cyvotoxic T lymphocytes, and these used for calculative immuncharany.
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Best Local S
Matches 9
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19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
                                                    Human; period (Drosophila) homologue 1; PER1; polymorphic variant; polymorphic site; genotyping; haplotyping; circadian rhythm regulation; single nucleotide polymorphism; SNP; gene; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis, prevention and treatment of cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 10
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(ROBE/)
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                                                                                                                     PER1 preferred oligonucleotide
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ROBERTS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunotherapy.
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                                                                                                                                                                                                                                                                                                                                                                         Conservative
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98US-0089997.
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Pred. No.
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                                                                                                                   primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 9
                                                                                                                                                                         Human; single nucleotide polymorphism; SNP; RANGAP1; haplotyping chromosome 22q13.2-q13.31; Ran GTPase activating protein 1; genocancer; irregular cell cycle associated disorder; primer; ss.
             17-APR-2000; 2000US-198072P
                                              17-APR-2001; 2001WO-US12455.
                                                                                                               WO200179240-A2
                                                                                                                                                                                                                                            Oligonucleotide #57 to detect human RANGAP1 gene polymorphisms.
                                                                                                                                                                                                                                                                              08-MAY-2002
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AAS19877 standard; DNA; 10
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                                                                                                                                              Homo sapiens
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9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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100.0%; Pr/
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ARESULT 58
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ID AAD266
AC AAD26
AC AADAC

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Best Local (
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                                                                                         New human endothelin 2 (EDN2) polymorphic variants and encoding useful in expressing EDN2 protein for screening candidate drugs treat diseases related to EDN2 activity -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; endothelin 2; EDN2; polymorphic site; PS; therapy; hyperte drug screening; cardiovascular disorder; renal insufficiency; ASO allele specific oligonucleotide; cerebroprotective; polymorphism;
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                                                                                                                                                                                                                    WPI; 2002-083075/11.
                                                                                                                                                                                                                                                                                 Kazemi A,
                                                                                                                                                                                                                                                                                                                                                                                                       19-MAY-2000; 2000US-205761P
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9; Conserv
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100.0%;
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Pred. No.
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o. 64;
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Claim 18; Page 15; 91pp; English

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Best Local S
Matches
has a tumour cell or site of angiogenesis, or if a treatment is effective in changing angiogenesis or changing a status of a set of target cells, comprising determining if a sample of the subject has an expression product of at least one marker gene. Also described is a compound capable of altering the expression or activity of Keratin 14, TIE 1, Salioadhesin or Siglec in a cell. Peripheral blood mononuclear cell (PBMC) expressed Keratin 14, TIE 1, Salioadhesin or Siglec, and kits containing them from the present invention can be used in a diagnostic method, particularly as
                                                                                                                                                                                                                                                                                                                                                     Determining presence of a tumor cell or angiogenesis, effectiveness of treatment, by detecting the presence is useful to detect and monitor treatment of Kaposi's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-JAN-2001; 2001EP-0200228
28-SEP-2001; 2001EP-0203703
28-SEP-2001; 2001US-325722P
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                                                                                                                                                                                                                                         The present invention describes a method for determining if an
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                                                                                                                                                                                                                                                                                                         e.g.
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                                                                               Sequence 11
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                                                                                                                   ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of skin. The present sequence is that of a human expressed sequence tag
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RESULT 62
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ABV64419/c
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Matches 9
 Human skin EST 2423
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RESULT 63
ABV6840/c
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AC ABV684
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                                                                                                                                          Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
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20-DEC-2001; 2001WO-EP15179

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RESULT 64
ABV68594/c
ID ABV685
XX ABV685
XX Human
XX Human;
KW Human;
KW immuno
KW Jeoria
XX WO2002
XX WO2002
XX HOMO 9
XX HOMO 9
XX PO 11-JUL
XX 20-DEC
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Disclosure; Page 202; 1345pp; German
                                                                                                                                                                                                                                                                                                                            Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
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e.g. skin ca
                                          In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
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         The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag
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 rosacea; melanoma; basal ce
skin. The present sequence
(EST) of the invention.
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immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis;
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Pred. No.
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RESULT 66
ABV71840/6
ID ABV718
XX ABV718
AC ABV71
AC ABV71
AC ABV71
XX 21-OC
DE Human
XX Human
XX Human
XX Human
XX Human
XX Human
XX Homo
XX W0200
DE 11-JU
XX 03-JE
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CCAGGCCAA 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              or pharmaceutical agents against
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                                                                                                                      DB 1;
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                                                                                                                                          Length 11;
                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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RESULT

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AAS19798/c
ID AAS197
XX AAS197
XX OB-MAY
XX ASO pr
XX Human;
KW Human;
KW Cancer
XW Cancer
XW Cancer
XW WO2001
XX WO2001
XX HOMO S
XX WO2001
XX WO2001
XX WO2001
XX WO2001
XX Genoty
PT Chew A
XX Genoty
PT determ
PT nuclec
PX Genoty
PT determ
PT nuclec
YX Genoty
PT determ
PT de
      AAV60744/c
ID AAV607
XX
AC AAV607
XX
OB-DEC
XX
DE Primer
XX
DE Primer
XX
PCR pr
KW PCR pr
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                                                                                                                                                                                                                                                                                                                                                                             RESULT 68
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SNPs) in the human Ran GTPase activating protein 1 (RANGAP1) gene located on chromosome 22q13.2-q13.31, and methods for haplotyping and/or genotyping the RANGAP1 gene. The methods of the invention make use of allele-specific oligonucleotides (ASOs) as probes and primers and/or primer-extension oligonucleotides for detecting the RANGAP1 gene polymorphisms. The polymucleotides and screened compounds are useful for treatment of diseases associated with RANGAP1 activity, such as cancer and other disorders associated with an irregular cell cycle. AAS19743-AAS19820 represent ASO primers for detecting human RANGAP1
PCR primer; amplification; yeast; UAS; upstream activating sequence; transcription terminator; cell cycle; Upstream Activation Sequence; UAS; promoter; phosphorylation; cyclin; cyclin-dependent kinase; CDK; vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; single nucleotide polymorphism; SNP; RANGAP1; haplotyping chromosome 22q13.2-q13.31; Ran GTPase activating protein 1; genoty cancer; irregular cell cycle associated disorder; ASO; primer; ss; allele-specific oligonucleotide.
                                                                                                                            Primer #2 for human
                                                                                                                                                                                           08-DEC-1998
                                                                                                                                                                                                                                                        AAV60744;
                                                                                                                                                                                                                                                                                                              AAV60744 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 15 BP; 3 A; 6 C; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene polymorphisms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to novel single nucleotide polymorphisms (SNPs) in the human Ran GTPase activating protein 1 (RANGAP1) gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 15; Page 15; 148pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        determining haplotype of individual, involves determining identity on nucleotide pair at specific polymorphic sites for two copies of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Genotyping human Ran GTPase activating protein 1 gene of individual for determining haplotype of individual, involves determining identity of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-075068/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-APR-2000; 2000US-198072P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ASO primer #56 to detect human RANGAP1 gene polymorphisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAS19798;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200179240-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENAISSANCE PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GTCCTGGCCCGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Choi JY,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                        (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Koshy
                                                                                                                            CDK4 codons 1-163
                                                                                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      44.0%;
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                                                                                                                                                                                                                                                                                                                 ВÞ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 8.8;
Pred. No. 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
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AAQ96791
XII AAQI
XXX
AC AAQI
XXX
DT 26-I
XXX
DE HIV
XXX
KW WO9
PN WO9
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                                                                                                                                                                                                                                                        RESULT 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    proteins. The method comprises administering a compound to a cell line, which contains a reporter gene linked to an Upstream Activation Sequence (UAS) and a promoter, where the UAS binds a transcription control factor (TCF) which is regulated through cyclin/cyclin-dependent kinase (CDK) phosphorylation. Also included in the construct is an effector gene providing a gene product to permit normal cyclin/CDK regulation of the TCF. Expression of the reporter gene is then analysed in the cell line, thereby determining whether the compound affects the normal regulation. The method can be used to identify inhibitors and activators of cyclins, CDKs, cyclin/CDK complexes, cyclin kinase activators of cyclins, CDKs, cyclin/CDK complexes, cyclin kinase can be used for stimulating growth of cells (as in wound healing), or regulating excessive cell growth and division (as in cancer therapy).
                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 10
                                                 Human immunodeficiency virus type
                                                                              HIV-1; AIDS; attenuation; vaccine; nef gene; avirulence; ss
                                                                                                                     HIV-1 NL4-3
                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Primers AAV60743-V60745 were used to PCR amplify codons 1-163 of the human cyclin-dependent kinase 4 (hCDK4). The amplified product was used to generate a fusion protein comprising part of the hCDK4 sequence linked to codons 154-302 of the yeast PHOB5 gene. The fusion protein is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Screening for agents that effect cell cycle regulatory proteins using a cell line that expresses a reporter gene in response to regulation through phosphorylation by a cyclin/CDK system
                                                                                                                                                      26-MAR-1996
                                                                                                                                                                                                                       AAQ96791 standard; DNA; 10 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linked to codons 154-302 of the yeast PHO85 gene. The fusion protein is used to screen for compounds that affect mammalian cell cycle regulatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-NOV-1996;
16-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 4; Page 75; 93pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bitter
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cyclin kinase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (BITT-) BITTECH INC
                                                                                                                                                                                                                                                                                                         12 GGCCTGGCCÁGÁ 1
                                                                                                                                                                                                                                                                                                                                         6 GGCCAGGCCAAA 17
                                                                                                                                                                                                                                                                                                                                                                            10;
                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                     nef
                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 4 A; 6 C; 5
                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       inhibitor; CKI; growth; wound healing; cancer therapy; ss.
                                                                                                                   gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      96US-0031968.
96US-0029127.
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                                                                                                                                                                                                                                                                                                                                                                                          44.0%;
                                                                                                                   nucleotide deletion 386.
                                                                                                                                                                                                                                                                                                                                                                                                                                           G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                      Score 8.8; DB 1
Pred. No. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                        Length 18
                                                                                                                                                                                                                                                                                                                                                                        Indels
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RESULT 70
AAT29327/c
ID AAT293
XX AAT293
XX AAT293
XX 25-MAR
DT 25-MAR
DT 28-JUN
XX 5'-pri
XW 5'-pri
KW 5'-pri
KW charac
KW identi
KW diseas
XX W09531
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XX W09531
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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14-FEB-1994;
21-FEB-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5'-primer; mammalian; G-protein coupled receptor; PCR primer kit; characterisation; biological samples; PCR amplification; indexing identification; cloning; analysis; genes; genome mapping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Attenuation of pathogenic HIV-1 strain NL4-3 involves deletion of 1 or more decanucleotides (AAQ96406-Q97018) from the nef gene and/or 1 or more decanucleotides (AAQ97019-097166) from the LTR region; the sequence of AAQ96406 corresponds to nucleotides 1-10 of the nef gene (AAQ96141). The resulting avirulent HIV strains are still capable of
                                                    WPI; 1996-010958/01.
                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAR-2003
28-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (AAQ96141). The resulting avirulent HIV strains are still capable inducing an immune response in humans, and enable the generation of therapeutic, diagnostic and targeting agents against HIV-1 infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New non-pathogenic HIV-1 strain carrying a deletion in its nef gene or LTR region - can be used in a vaccine to inhibit/reduce productive infection in an individual by a pathogenic strain
                                                                                                       Lopeznieto CE,
                                                                                                                                                                                                           16-MAY-1994;
                                                                                                                                                                                                                                                        12-MAY-1995;
                                                                                                                                                                                                                                                                                                             23-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  disease diagnosis; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5'-primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT29327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT29327 standard; DNA; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 10 BP; 2 A; 3 C; 5 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 13; Page 193; 301pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1995-293115/38
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-AUG-1995
                                                                                                                                                         (BGHM ) BRIGHAM & WOMENS HOSPITAL.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3 CCGGGCCAGG 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for mammalian G-protein coupled receptor coding sequences.
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(first en
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94AU-0003864.
94AU-0004002.
                                                                                                    Nigam
                                                                                                                                                                                                           94US-0242887
                                                                                                                                                                                                                                                              95WO-US06032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cloning;
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90.0%;
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BURNET CENT MEDICAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
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                                                                                                       SK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         analysis; genes; genome mapping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 8.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pred.
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Characterisation of nucleotide sequences using primer pairs -

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PCR

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RESULT 71
AAV50125/C
ID AAV50
XX AAV50125/C
XX AAV50
AC AAV50
XX Yeast
YA (UYJU
XX YEAST
DR WPI;
XX Yeast
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Best Local S
Matches 9
             Yeast transcriptome is encoded by a DNA molecule comprising a yeast gene involved in cell cycle progression selected from the group of nonannotated ORF (NORF) genes SAGE (serial analysis gene expression) tags for highly expressed genes and NORF genes are given in AAV50051 to AAV50345. The present invention describes: (1) a method of using yeast genes to modulate the cell cycle which comprises administering to a cell an isolated DNA molecule comprising a yeast gene which is involved in cell cycle progression selected from differentially expressed genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The 5'-primers AAT29262-382, and the complementary 3'-primers derived from them, which target mammalian G-protein coupled receptor coding sequences, together comprise a PCR primer kit. The kit is used in a new method for the characterisation of nucleic acid sequences obtd. from mammalian biological samples, which comprises PCR amplification and indexing of the prods. w.r.t the primer pair that hybridised to its delineating subsequences. The method may be used in the identification, cloning and analysis of genes, e.g. in genome mapping, and disease diagnosis.
                                                                                                                                                                                                                                                                         Claim
                                                                                                                                                                                                                                                                                                                                                 Yeast transcriptome - useful for screening antifungal agents, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yeast; Saccharomyces cerevisiae; transcriptome; cell cycle; regulation; eukaryotic cell; antifungal; SAGE tag; gene expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-427943/36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-JAN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UYJO ) UNIV JOHNS HOPKINS SCHOOL MEDICINE
cycle progression selected
E tags given in AAV50051 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tag
                                                                                                                                                                                                                                                                      1; Page 23;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      for NORF gene locus NORF15.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gene expression;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 8.4;
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AAV50345);
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                                                                                                                                                                                                                                                                                                                                               modulating eukaryotic cell, for identifying genes in cell
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RESULT 72
AX54917
ID AXX54
AC AAX54
AC AAX54
AC AAX54
AX ACCEBP
XX Antis
KW Inpai
KW inpai
KW pulmo
KW cohron
KW cohron
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KW pulmo
KW resource
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  candidate antifungal drugs which comprises contacting a test substance with a yeast cell and monitoring expression of a yeast gene which is involved in cell cycle progression; (3) a method of identifying human genes which are involved in cell cycle progression which comprises hybridizing a probe comprising at least 10 contiguous nucleotides of a yeast gene which is differentially expressed between at least 2 phases selected from the log phase, the S phase and the G2/M phase; and (4) a probe for ascertaining the phase in the cell cycle, where the probe comprises at least 14 contiguous nucleotides of a NORF gene (SAGE tags given in AAV50051 to AAV50345), or as an array of probes on a solid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer; hepatic metastasis
                                                                                                                                                                                                                                                                                                             09-JUN-1998;
17-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                            25-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         acute asthma; allergy; asthma; impeded respiration;
respiratory distress syndrome; pain; cystic fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         pulmonary vasoconstriction; inflammation; allergic rhinitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C/EBP-beta antisense oligonucleotide fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-JUL-1999
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                                                                                                                                                                            New antisense oligonucleotides used in
                                                                                                                                                                                                                                                                                                                                                            17-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                            WO9913886-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        impaired respiration; inflammation; lung disease;
                                                                                                                                                                                                                                                                             (UYEC-)
                                                                                                                                                                                                              1999-229400/19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hypertension; pulmonary vasoconstriction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP;
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                                                                                                                                                                                                                                                                              EAST CAROLINA
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97US-0059160
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             88.
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90.0%;
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Pred.
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                                                                                                                                                                              treatment
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                                                                                                                                                                              pulmonary
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The specification describes antisense oligonucleotides (AAX52869-X55271) directed against at least 2 mRNAs selected from target genes, coding and non-coding regions of RNAs corresponding to target genes, gene initiation codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'-end and the juxta-section between coding and non-coding regions and all segments of RNAs encoding proteins associated with one or more diseases, conditions or mixtures. The antisense oligonucleotides

Disclosure; Page 68; 120pp; English

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RESULT 73
AAF20486
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                                                                    Low adenosine (A) content antisense oligonucleotides which do not trigger adenosine receptors during metabolism, useful e.g. for tr cancers and respiratory obstructions -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory discress syndrome; pain; cystic fibrosis; allergic rhinitis; pulmonary hypertension; emphysema; pulmonary transplantation rejection; chronic obstructive pulmonary disease; pulmonary infection; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Low adenosine antisense oligonucleotide; phosphorothioate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic; respiratory obstruction; pulmonary obstruction; impeded respiration; respiratory obstruction; pulmonary obstruction; impeded respiration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  may be derived from sequences AAX55272-74. These multiple target oligonucleotides (specifically AAX55180-271) can be used for the antisense treatment of diseases and conditions. Typical diseases conditions are those associated with impaired respiration and
Claim 14; Page 262; 1592pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-MAR-2000; 2000WO-US08020.
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                                                                                                                                                                                                          WPI; 2000-679539/66
                                                                                                                                                                                                                                                                                                                                                   (UYEC-) UNIV EAST CAROLINA. (NYCE/) NYCE J W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human C/EBP polynucleotide fragment #2053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAF20486 standard; DNA; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 10 BP; 0 A; 4 C; 6 G; 0 U; 0 other;
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9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CGGGCCAGGC 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CGGGCCGGGC 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                        99US-0127958
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Pred. No. 87;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB
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                                                                                                     for treating
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The present invention describes low adenosine (A) content antisense oligonucleotides and compositions (I) comprising them. In the antisense oligonucleotides the A is replaced by a 'Universal' or alternative base

Claim 1;

Page 9;

95pp; Japanese

autoimmune disease and tumors

The present invention describes a group of genes consisting of 100 genes

bronchodilator,

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AAC73935/c
ID AAC73935/c
ID AAC7399
XX AC7399
XX O2-FEB
XX Human;
XW Human;
XW autoim
XX HOMO S
PN WO2000
XX HOMO S
PN WO1-APR
XX WPI; 2
XX Groups
PT Autoin
XX Claim
XX Claim
XX Claim
XX Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. CC The antisense oligonuclectides and (I) can be used to down-regulate the CC expression and or activity of target polypeptides associated with CC lung/respiratory disorders and malignancies, such as stimulating and CC immunoglobulins and antibodies, antibody receptors, cytokines and CC chemokines, endogenously produced specific and non-specific enzymes, CC binding proteins, adhesion molecules and their receptors, cytokine and CC chemokines receptors, adenosine receptors, bradykinin receptors, central CC nervous system (CNS) and peripheral nervous and non-nervous system peptide receptors, conditions and peripheral nervous and non-nervous system peptide concluding respiratory obstruction and malignancy associated proteins. The CC including respiratory obstruction (especially pulmonary obstruction and/or bronchoconstriction) and/or lung inflammation, allergy(res) and/or surfactant hypoproduction which are associated with a disease or condition selected from pulmonary vasoconstriction, inflammation, callergies, asthma, impeded respiration, respiratory distress syndrome (CC plumonary transplantation rejection, pulmonary inflammation, bronchitis, and constructive pulmonary disease (COPD), pain, cystic fibrosis (CF), allergic rhinitis (AR), pulmonary conservation, emphysema, chronic obstructive pulmonary disease (COPD), concert. AAF18434 to AAF21543 represent human polymucleotide conservation of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 74
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
                                                                                            Groups of genes expressed in human dendritic cells lesser extent than in monocytes for investigation a
                                                                                                                                                      WPI; 2000-619172/59.
                                                                                                                                                                                          Hashimoto S,
                                                                                                                                                                                                                                                                    01-APR-1999;
                                                                                                                                                                                                                                                                                                       30-MAR-2000; 2000WO-JP02019
                                                                                                                                                                                                                                                                                                                                                                                  WO200060074-A1
                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; dendritic cell; monocyte; immune system; diagnosis; cancer;
autoimmune disease; tumour; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human dendritic cell cDNA base sequence oligonucleotide #22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-FEB-2001
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9; Conserv
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90.0%;
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                                                                                                                                                                                          Suzuki T
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Pred. No. 8
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diagnosis
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RESULT 75
AAC74189/c
ID AAC741
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XX AAC741
XX AAC741
XX D2-FEB
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Matches
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The present invention describes a group of genes consisting of 100 genes which are highly expressed in human dendritic cells; a group of genes which are expressed at a higher frequency in human dendritic cells than in human monocytes; and a group of genes which are expressed at lower frequency in human dendritic cells than in human monocytes. Each group of genes are characterised in that CDNAs of these genes respectively have the base sequences of SEQ ID NO:101 to 100 (AAC74013) and SEQ ID NO:201 to 300 (AAC74014 to AAC74013), each is continuous with the base sequence 5'-CATG-3' located most closely to the poly-A region. The sequence 5'-CATG-3' located most closely to the poly-A region. The sequence 5'-CATG-3' located most closely to the poly-A region. The sequence so the used for the investigation of the role and mechanism of the involvement of dendritic cells in the immune system and for the study and diagnosis of diseases in which dendritic cells play a significant role,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Groups of genes expressed in human lesser extent than in monocytes for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-619172/59
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; dendritic cell; monocyte; immune system; diagnosis; cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disease and tumors
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                                                    The present invention describes a new composition comprising an antisense oligonucleotide (ON) with low adenosine (up to 15%), which CC targets nucleic acids involved in bronchoconstriction, allergies, and/or inflammation. The ON can have antiinflammatory, antiallergie, inflammation. The ON can have antiinflammatory, antiallergie, are conserved airways, including lung disease activities. The compositions are compared airways, including lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating ce.g. ischaemic conditions, pulmonary vasoconstriction, allergies, casthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, emphysema, chronic obstructive conditions, pulmotary disease (COPD), and cancers such as leukaemias, lymphomas, concerniomas, and cancers which may metastasise to the lungs, including conditions, and cancers which may metastasise to the adenosine content of the ONs reduces side effects. The reduction of the adenosine content of the case of decayadenosine which activates adenosine receptors causing the proposed the content of the content of the case of decayadenosine which activates adenosine receptors causing the proposed the content of the content of the case of decayadenosine which activates adenosine receptors causing the proposed the content of the content of the case of decayadenosine which activates adenosine receptors causing the proposed to the content of the content of the case of decayadenosine which activates adenosine receptors causing the proposed the content of the content of the content of the content of the case of decayadenosine which activates adenosine receptors causing the proposed the content of th
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstriction, inflammation, allergies, asthma, hypertension,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      respiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; Cancer; leukaemia; lymphoma; carcinoma; metastasis; ss.
bronchoconstriction and inflammation. AAA32313 to AAA35312 nucleotide sequences given in the sequence listing from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           bronchitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-205971/18.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human adenosine receptor related polynucleotide SEQ ID NO: 2053
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 523;
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90.0%;
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                                                                                                                                                                                                                                                                                                                                              SAGE tag; serial analysis of gene expression; antigen-presenting ce APC; monocyte-derived dendritic cell; differential gene expression; immunostimulatory cofactor; costimulatory factor; CTL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention, which correspond to SEQ ID NO:1 to 2815, and then the last 185 sequences are also called SEQ ID NO:1 to 185, but the sequences differ from the previously named sequences. SEQ ID NO:11 to 1880 (AAA32323 to AAA33992) are specifically claimed ONs from the present invention. N.B. Sequences given in the disclosure of the present invention do not match up with their corresponding SEQ ID NO: sequences given in the sequence listing.
                                                                                                                  19-JUN-
                                                                                                                                                                                                                                                                                                                                                                                Human dendritic cell SAGE tag,
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                                                                                                                                                                                                                                                                                                                                       immunotherapy; anticancer; ss
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RESULT 78 AAZ78898

AAZ78898;

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(first entry)

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Query Match Best Local S Matches 9

Similarity

90 90

0.0%

Score 8.4; I Pred. No. 87

DB

Length 10 Indels

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Conservative

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Mismatches

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Gaps

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Sequence 10

BP; 0 A; 3 C; 5 G; 2 T; 0 other;

Human dendritic cell SAGE tag,

SEQ ID NO:1326

SAGE tag; serial analysis of gene expression; antigen-presenting ce APC; monocyte-derived dendritic cell; differential gene expression; immunostimulatory cofactor; costimulatory factor; CTL;

cell;

Homo sapiens

T-lymphocyte;

tumour antigen;

immunotherapy;

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context transcripts correspond to novel genes in diversity collections (APC)-associated costimulatory factors play an important role in the complex cells. Tumour antigen presentation via the MHC (major histocompatibility complex) and subsequent recognition by T-cell receptors is alone complex) and subsequent recognition by T-cell receptors is alone complex) and subsequent recognition by T-cell receptors is alone complex. Inductivate a robust cytotoxic immune response that can complex is the tumour cells, immunostimulatory cofactors also being required complex is dentified using the cytotoxic T-lymphocytes (CTLs). Nucleic acid complex is dentified using the cytotoxic T-lymphocytes (CTLs). Nucleic acid complex is a tumour antigen; to modulate represent ally expressed genes in complex is a tumour antigen; to modulate an immune response, particularly against a tumour antigen; to modulate the genotype of an APC; to screen complex is an antigen; to modulate the genotype of an APC; to screen complex is an antigen; to modulate the genotype of an APC; to screen complex is an antigen; to the monocyte lineage. Cells containing composis, prognosis and monitoring of diseases related to abnormal compression of these genes, or of their encoded proteins, can be used to identify cells as belonging to the monocyte lineage. Cells containing them are used in active immunotherapy (or to stimulate containing them are used in gene therapy. Co-administration of cumour antigens and APC-associated costimulatory factors ensures adequate antigen presentation to endogenous APCs and upregulates the APCs for the containment of immune effector cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  with monocytes. Some of the transcripts correspond to known genes or ESTs (expressed sequence tags) which were previously unknown to be preferentially or differentially expressed in dendritic cells, while
                                                     recruitment of immune effector cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        differentially expressed in monocyte-derived dendritic cells compared with monocytes. Some of the transcripts correspond to known genes or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequences AAZ77573-Z79709 represent SAGE (serial analysis of expression) tags used to identify mRNA transcripts encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Isolated polynucleotides differentially expressed in antigen-presenting cells, useful in gene vaccines against cancer - \,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ROBE/) ROBERTS B L. (SHAN/) SHANKARA S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mmunostimulatory cofactor proteins which are preferentially or
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18-JUN-1999;

99WO-US13800

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Sequences AAZ77573-Z79709 represent SAGE (serial analysis of gene cxpression) tags used to identify mRNA transcripts encoding immunostimulatory cofactor proteins which are preferentially or CC differentially expressed in monocyte-derived dendritic cells compared with monocytes. Some of the transcripts correspond to known genes or CC ESTs (expressed sequence tags) which were previously unknown to be preferentially or differentially expressed in dendritic cells, while CC (APC) associated costimulatory factors play an important role in the CC (apc) associated costimulatory factors play an important role in the CC complex) and subsequent recognition by T-cell receptors is alone complex) and subsequent recognition by T-cell receptors is alone constituted to activate a robust cytotoxic immune response that can complex and subsequent recognition by T-cell receptors is alone constituted a crivate a robust cytotoxic immune response that can constituted using the SAGE tags have several potential uses. They may be used in vaccines to induce an immune response, particularly against a tumour antigen; to modulate the genotype of an APC; to screen for agents that modulate expression of differentially expressed genes in CC an APC; and as hybridisation probes/amplification primers for the cc an APC; and as hybridisation probes/amplification primers for the cc differentially expressed genes. Detection of the dendritic cell containing to identify cells as belonging to the monocyte lineage. Cells containing
           Isolated polynucleotides differentially expressed in antigen-presenting cells, useful in gene vaccines against cancer -
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RESULT 79
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       AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells).

AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour cissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisaction/amplification reactions. Compounds that modulate expression of the transcripts are
                                                                                                                                                                         Claim 1; Page 79; 219pp; English.
                                                                                                                                                                                                        prevention and
                                                                                                                                                                                                                    Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis,
                                                                                                                                                                                                                                                                                             Roberts BL,
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               AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic.
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                                                                                                                                                                                                                                                                                                         Isolated polynucleotides differentially and non-metastatic breast cancer cells, prevention and treatment of cancer -
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  or hybridisation/amplification
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AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells).

AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis,

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AAZ80767 to AAZ83941 represent tags transcripts that are preferentially tumour tissue (i.e. are upregulated AAZ83942 to AAZ86677 represent tags that are preferentially transcribed
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                                                                                          Isolated polynucleotides differentially and non-metastatic breast cancer cells, prevention and treatment of cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; metastatic breast tumour tissue; breast cancer; tag; pr non-metastatic breast tumour tissue; gene therapy; anticancer; antimetastatic; vaccine; diagnosis; ss.
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AAZ80767 to AAZ83941 represent tags transcripts that are preferentially tumour tissue (i.e. are upregulated

corresponding to distinct transcribed in the metastatic in metastatic breast tumour co

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) SHANKARA S.
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antimetastatic; vaccine; diagnosis; ss.
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Claim 1; Page 148; 219pp; English

prevention and

treatment

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RESULT 86
AAZ84369/c
ID AAZ84369 standard; DNA; 10 BI
XX
AC AAZ84369;
XX
AC AAZ84369;
XX
DT 07-APR-2000 (first entry)
XX
DE Metastatic breast tumour cell
XX
DE Metastatic breast tumour cell
XX
Human; metastatic; vaccine; diag
XX
Homo sapiens.
XX
AC AAZ84369;
XX
Homo sapiens.
XX
PN W09965928-A2.
XX
PN W099
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Best Local S
Matches
            Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis, prevention and treatment of cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; metastatic breast tumour tissue; breast cancer; tag; primer;
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98US-0089997.
98US-0090039.
98US-0090040.
98US-0090041.
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Pred. No. 8
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ARESULT 87
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Best Local
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19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
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Isolated polynucleotides differentially expressed between metastatic
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(ROBE/)
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antimetastatic; vaccine;
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ROBERTS B L.
SHANKARA S.
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ne; diagnosis; ss.
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Pred. No. 87;
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Matches 9
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                                                                                                                                                                                                                                                                                                                                            Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer; antimetastatic; vaccine; diagnosis; ss.
WPI; 2000-106079/09
                             Roberts BL,
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19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                  Homo sapiens
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prevention and treatment of cancer -
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19-JUN-1998;
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                                                             (SHAN/)
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                                                                           GENZYME CORP. ROBERTS B L.
                                                             SHANKARA S.
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                             Shankara
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Pred. No. 87;
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reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in select cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be use of expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for
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AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification
Sequence 10 BP; 0 A; 3 C; 5 G; 2 T; 0 other;
                                                  adoptive immunotherapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   prevention and treatment of cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Isolated polynucleotides differentially and non-metastatic breast cancer cells,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 171; 219pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          expressed between metastatic useful for diagnosis,
                                                                                                                                                                                                                                                                                            in selected
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Query Match
Best Local S
Matches 9
10 GCCAGGCCCA 1
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                                Similarity
9; Conserv
               GCCAGGCCAA 16
                                 Conservative
                                         42.0%;
                                0; Mismatches
                                         Score 8.4; D
Pred. No. 87;
                                                 DB 1;
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                                                  Length 10
                                 Indels
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                                Gaps
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RESULT 89
AAZ85815/c
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
Roberts BL,
                           (GENZ )
(ROBE/)
                                                                                                                            23-DEC-1999
                                                                                                                                            WO9965928-A2
                                                                                                                                                             Homo sapiens
                                                                                                                                                                               antimetastatic; vaccine; diagnosis; ss.
                                                                                                                                                                                      non-metastatic breast tumour tissue; gene therapy; anticancer;
                                                                                                                                                                                                Human; metastatic breast tumour tissue; breast cancer; tag;
                                                                                                                                                                                                                 Metastatic breast tumour cell downregulated transcript tag #5049
                                                                                                                                                                                                                                    07-APR-2000
                                                                                                                                                                                                                                                                      AAZ85815
                                                                               19-JUN-1998;
19-JUN-1998;
                                                                                                          18-JUN-1999;
                  (SHAN/)
                         GENZYME CORP
ROBERTS B L.
                  SHANKARA S
                                                                                                                                                                                                                                                                      standard; DNA; 10
Shankara
                                                                                                                                                                                                                                   (first entry)
                                                                                                          99WO-US13647
                                                      98US-0090041
                                                              98US-0090040
                                                                      98US-0090039
                                                                               98US-0089997
                                                                                        98US-0089853
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RESULT 90
AAZ85951
AAZ85951
XX

XX

AC AAZ85
AC AAZ85
AC Meta
XX

DE Meta
XX

Huma
KW Huma
KW ann:
XX

OS Hom
YX

YX

PN WO9
YX

PF 18:
XX

PF 18:
YX

PR 19
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CC AAR33942 to AAR36677 represent tags corresponding to distinct transcripts come that are preferentially transcribed in the primary or non-metastatic breast tumour cells).

CC that are preferentially transcribed in the primary or non-metastatic come that the primary or non-metastatic breast tumour composition of the transcripts can be used for diagnosis, prognosis, composition of breast cancer, particularly where metastatic Dreast tumour compositions. Compounds that modulate expression of the transcripts are concertable useful for treatment of (metastatic) breast cancer, while compounds that modulate expression of the transcripts are concertable useful for treatment of (metastatic) breast cancer, while compounds that modulate expression of the transcripts are concertable, particularly an antigenes (also ribozymes or antisense concertable), particularly an antigenes (also ribozymes or antisense or cell-based vaccines, Polypetides encoded by the transcripts are also concertable of the polypetides or as the polypetides of as the polypetides of as the polypetides can be used to expand and isolate populations of educated, antigen-specific immune cancer cells, e.g. cytotoxic T lymphocytes, and these used for
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Best Local
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                                                                                                                                  19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis, prevention and treatment of cancer -
      (GENZ )
(ROBE/)
(SHAN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  non-metastatic breast tumour tissue; gene therapy; anticancer; antimetastatic; vaccine; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; metastatic breast tumour tissue; breast cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-APR-2000
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                                                                                                                                                                                                                                19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                 18-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Metastatic breast tumour cell downregulated transcript tag #5185.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ85951 standard; DNA; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2000-106079/09
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      GENZYME CORP.
ROBERTS B L.
SHANKARA S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunotherapy.
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                                                                                                                               98US-0089853.
98US-0089997.
98US-0090039.
98US-0090040.
98US-0090041.
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90.0%;
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reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines, for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for
                                                                                                                                                                                                                                                                                                                                             AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnossis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic. Diagnossis is by standard immunoassays or hybridisation/amplification
Sequence 10
                                                 adoptive immunotherapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 196; 219pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Roberts BL,
BP; 1 A; 5 C; 4
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G; 0 U; 0 other;
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                           Query Match
Best Local S
Matches 9
                          l Similarity
9; Conserv
             ហ
GGGCCAGCCC
             GGGCCAGGCC
                           Conservative
             14
10
                                 42.0%;
90.0%;
                                  Pred. No. 87
                           Mismatches
                                         DB 1;
                           1;
                                       Length 10,
                           Indels
                           0
                           Gaps
                           0
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AAZ86053/c
II AAZ860)
XX
AC AAZ860)
XX
DT 07-APR
XX Metast
XX
Human;
KW Human;
KW non-me
KW antime
XX Homo s
XX
PN WO9965
XX HOMO S
XX
PN WO9965
XX 19-JUN
PR 19-JUN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9965928-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    non-metastatic breast tur
antimetastatic; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Metastatic breast tumour cell downregulated transcript
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        metastatic breast tumour tissue; breast cancer; tag;
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ne; diagnosis; ss.
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tag

#5287.

19-JUN-1998; 19-JUN-1998; 19-JUN-1998; 19-JUN-1998;

98US-0089853. 98US-0089997. 98US-0090039. 98US-0090040.

(GENZ ) GENZYME CORP

18-JUN-1999;

99WO-US13647

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RESULT 92
AA167394/c
ID AA1673
XX AA1673
XX AA1673
XX I1-FEB
XX Human
XX FK506-
KW FK506-
KW immuno
XX Homo s
XX W02001
XX W2001
XX PF 26-MAR
XX 26-MAR
XX AAASTA
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metasta Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for trastment of (metastatic) breast cancer, while
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour
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               Anastasio
                                                                                                                26-MAR-2001; 2001WO-US09718
                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                           FK506-binding protein 8; FKBP8; haplotyping; polymorphism; cancer;
                                                                                                                                                                                                                                                                                             Human FKBP8 gene polymorphism detecting primer
                                                                                                                                                                                                                                                                                                                                 11-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 10 BP; 0 A; 4 C; 4
                                                                                24-MAR-2000; 2000US-192125P.
                                                                                                                                                                                WO200172965-A2
                                                                                                                                                                                                                                                                                                                                                                AAI67394;
                                                                                                                                                                                                                                                                                                                                                                                               AAI67394 standard;
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                                                 (GENA-) GENAISSANCE PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             promoters from the transcripts are used to direct expression,
                                                                                                                                                                                                                                               mmunosuppression;
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                 Bentivegna SC,
                                                                                                                                                                                                                                               human; primer;
                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
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90.0%;
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                 Choi JY,
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                 Kliem
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               SE,
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ABA06204/c
ID ABA062
XX ABA062
AC ABA062
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AC Human
ID Human;
XX Human;
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CC Group
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                                                                                      The invention relates to a human normal hepatocyte expression gene group comprising 200 genes in the human normal hepatocyte. The cDNA of each gene comprises one of 200 fully defined nucleotide sequences as given in the specification. The gene group and the cDN corresponding to each of the genes in the group are useful in the diagnosis and treatment of human hepatopathy. The present sequence is a cDNA corresponding to a gene expressed by normal human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to haplotyping the FK506-binding protein 8 (38kD) (FKBP8) gene in an individual. The method involves determining the identity of the nucleotide pair at one or more polymorphic sites selected from P1 to P26 (described in the specification). The invention is useful to improve the efficiency and reliability of several steps in the discovery and development of drugs for treating diseases associated with FKBP8 activity, for example immunosuppression and cancer. Sequences
                                                                                                                                                                                                                                                                                                                                   Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-629566/73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-JAN-2000; 2000JP-0023170.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; hepatocyte; gene expression; hepatopathy; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human normal hepatocyte
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-JAN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 10 BP; 0 A; 2 C; 5 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                polymorphisms by primer extension techniques.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 16; Page 15; 98pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                genotyping that gene in individual and to design new the associated disease such as immunosuppression and cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New haplotypes of the FK506-binding protein 8 gene, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (KAGA-) KAGAKU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-626261/72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                                                                        normal hepatocyte expression
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity 9; Conserv
                                                                                                                                                                                                                                                                                                                                Page 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
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                                                                                                                                                                                                                                                                                                                                   26pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
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90.0%;
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                                                                                                                                                                                                                                                                                                                                                                                        gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene cDNA, SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  No. 87;
                                                                                                                                                                                                                                                                                                                                                                                        group
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for detecting FKBP8 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 10;
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Query Match Best Local Similarity

42.0**%**; 90.0**%**;

Score 8.4; D Pred. No. 87;

DB 1;

Length 10;

Sequence 10

BP; 1 A; 3 C; 4 G; 2 T; 0 other;

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RESULT 94
AAH76342/c
ID AAH763
RESULT 95
AAH76345
ID AAH76
XX
AC AAH76
XX
DT 29-OC
XX
DE Z. ma
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                                                                                                                                      Query Match
Best Local S
Matches 9
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                                                                                                                                                                                                           The invention provides a male tissue-preferred regulatory region (I) comprising nucleotide sequences essential for initiating transcription of the MS45 gene. A method of mediating male fertility in a plant is provided that involves introducing an expression vector comprising a promoter operably linked to (I) into a plant where the exogenous gene impacts male fertility of the plant and (I) controls expression of the exogenous gene. A method of producing hybrid seeds is also provided. Sequences AAH763A1-355 represent a series of 5' deletions in the MS45
                                                                                                                                                                                                                                                                                                               A male tissue-preferred regulatory region comprising nucleotide sequences essential for initiating transcription of the MS45 geuseful for mediating fertility in a male plant -
                                      AAH76345;
                                                        AAH76345 standard;
                                                                                                                                                                          Sequence 10 BP;
                                                                                                                                                                                                                                                                                              Example 5; Fig 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mays Ms45 promoter deletion mutant fragment LS02.
                    29-OCT-2001
                                                                                                                                                                                                    promoter region, used for
                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-514772/56
                                                                                                                                                                                                                                                                                                                                                                                                                           13-FEB-2001; 2001WO-US04527.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Zea mays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hybrid seed;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAH76342;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAH76342 standard; DNA; 10 BP
                                                                                                                                                                                                                                                                                                                                                                                        (PION-) PIONEER HI-BRED INT INC
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mays Ms45 promoter deletion mutant fragment LS05
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                                                                                                                    4 CGGGCCAGGC 13
                                                                                                                                      Similarity 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tissue; regulatory region; transcription; male fertility;
                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   promoter;
                   (first entry)
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                                                        DNA; 10 BP
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                                                                                                                                                                                                    determining the essential region of MS45
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                                                                                                                                                       Score 8.4; DB 1;
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AAH76348/c
ID AAH763
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AC AAH763
AC AAH763
AC Z- may
XX
DF Z- may
XX
KW Ms45;
KW Mybrid
XX
Cea ma
XX
W02001
XX
PD 23-AUG
PD 23-FEB
XX
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XX
PION-
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13-FEB-2001; 2001WO-US04527

(PION-) PIONEER HI-BRED INT INC 15-FEB-2000; 2000US-0504487 23-AUG-2001 WO200160997-A2 hybrid

seed;

promoter;

29-OCT-2001

(first

entry)

Ms45; male tissue; regulatory region; transcription; mays Ms45 promoter deletion mutant fragment LS08.

male fertility;

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RESULT 96
                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                               Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                  provided that involves introducing an expression vector comprising a promoter operably linked to (I) into a plant where the exogenous gene impacts male fertility of the plant and (I) controls expression of the exogenous gene. A method of producing hybrid seeds is also provided. Sequences AAH76341-355 represent a series of 5' deletions in the Ms45 promoter region, used for determining the essential region of MS45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A male tissue-preferred regulatory region comprising nucleotide sequences essential for initiating transcription of the MS45 gene
AAH76348
                                    AAH76348 standard;
                                                                                                                                                                                                                                                                                                        Sequence 10 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           comprising nucleotide sequences essential for initiating transcription of the MS45 gene. A method of mediating male fertility in a plant is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention provides a male tissue-preferred regulatory region (I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 5; Fig 8; 50pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ms45; male tissue; regulatory region; transcription; male fertility; hybrid seed; promoter; ss.
                                                                                                                                                                                                                                                                                                                                                     promoter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-FEB-2000; 2000US-0504487
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                                                                                                                                                                                                                       Similarity
9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mediating fertility in a male plant
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                                      DNA;
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                                      ВP
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                                                                                                                                                                                                                                                                                                          G; 0 U; 0 other;
                                                                                                                                                                                                                                            Score 8.4; D
Pred. No. 87;
                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                Length 10;
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RESULT 97
AAH63580/c
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Best Local S
Matches 9
The present invention describes a method of identifying the type of cell in a sample, involving determining which of the sequences ARH63161-ARH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention provides a male tissue-preferred regulatory region (I) comprising nucleotide sequences essential for initiating transcription of the MS45 gene. A method of mediating male fertility in a plant is provided that involves introducing an expression vector comprising a promoter operably linked to (I) into a plant where the exogenous gene impacts male fertility of the plant and (I) controls expression of the exogenous gene. A method of producing hybrid seeds is also provided. Sequences AAH76341-355 represent a series of 5' deletions in the MS45 promoter region, used for determining the essential region of MS45
                                                                                                              New isolated polynucleotides, type, such as cancer cell, con particular cell types -
                                                                                                                                                                                                                                                                                                                                                                  Human; transcriptome; gene expression pattern; cancer; cancer diagnosis; cell specific gene expression; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A male tissue-preferred regulatory region comprising sequences essential for initiating transcription of tuseful for mediating fertility in a male plant -
                                                                                                                                                                                         Velculescu
                                                                                                                                                                                                                                                                                            31-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                      Human ubiquitously expressed transcriptome sequence SEQ ID NO: 420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 10
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                                                                                                                                                                                                                                                                  21-NOV-2000; 2000WO-US31922
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                                                                                                                                                                                                                                                                                                                                            Homo
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                                                                                                                                                                                                                                                                                                                                            sapiens
                                                                                       13; Page 48; 94pp;
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9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CDNA;
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AAH61161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce
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The present invention describes a method of identifying the type of cell in a sample, involving determining which for the sequences AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce
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particular cell types -
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RESULT 102
AAH64389/c
ID AAH643
XX AAH643
XX AAH643
XX AAH643
XX Human
XX Human
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XX Homo s
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  RESULT 103
AAH64390/c
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AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce
AAH64390 standard; cDNA;
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Pred. No. 87;
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21-NOV-2000; 2000WO-US31922.

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RESULT 104
AAH64391/c
ID AAH643
XX AAH643
XX AAH643
XX CONSEP
XX Human
XX Human;
XW Human;
XW Cancer
XX Cancer
XX WO2001
XX WO2001
XX XX
PD 31-MAY
XX
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                                                                                                                                                                              Human; cancer
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9; Conserv
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specific gene expression; ss.
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RESULT 105
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XX AAH643
XX Human
XX Human
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XX Homo 8
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The present invention describes a method of identifying the type of cell in a sample, involving determining which of the sequences AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously
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                                                                                                                                                                                                                                                                                                                                                     Velculescu VE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-NOV-2000; 2000WO-US31922
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                                                                                                                                       Page 67; 94pp; English.
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RESULT 106
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XX AAH643
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RESULT 107 AAH64394/c

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RESULT 108
AAH33882/C
ID AAH328
XX AAH328
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DT 13-AUG
DT 13-AUG
CXX Human;
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                          21-MAR-2001
                                                                                                             Human; LPS; lipopolysaccharide; monocyte
                                                                                                                                                                 13-AUG-2001
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28-APR-2000; 2000JP-0131079.
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                                                                                                  expressed sequence
                                                                                                                                    LPS activated human monocyte expression gene cDNA tag SEQ:255
                                                                                                                                                                                          AAH32882;
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RESULT 109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes an lipopolysaccharide (LPS) activated human monocyte expression gene group consisting of the high-ranking 50 genes of the high-stanking 50 genes of the high-stanking 50 genes of the high-stanking 50 genes expressed by LPS in which the cDNA of each gene has the base sequence of (AAH32628 to AAH32677) continuous to the base sequence 5'-CATG-3' nearest to the polyA region. The gene group is useful for the development of new means for the diagnosis and the treatment of various human diseases in which human monocyte plays an important role. AAH32628 to AAH32943 represent specifically claimed LPS activated human monocyte expression gene cDNA tags from the present invention. AAH32944 represents an LPS activated human monocyte expression gene cDNA sequence encoding AAB98009, which are given in the exemplification of the present invention.
                                                                                                                         New isolated polynucleotide comprising a polymorphic variant solute carrier family 6 neurotransmitter transporter, seroto 4 gene for identifying drugs for treating disorders related
                                                   Disclosure;
                                                                                                         expression of the protein
                                                                                                                                                                                                                                                                                        Denton
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                                                Page
                                                                                                                                                                                                                                                                                   Duda A,
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The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M2) for identifying human genes which are involved in cell cycle progression comprising contacting human DNA with a probe which comprises as in M1, and (4) a method (M4) for identifying a candidate drug as a comber of a class of drugs having a characteristic effect on gene
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The invention is used in producing a recombinant organism that can be used to express SLC6A4 for protein structure analysis and binding studies. A composition comprising a genotyping oligonucleotide is used to detect a polymorphism in the SLC6A4 gene.
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The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by alleast 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression varies as in M1, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M2) for identifying hyper capacity and in call order.

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ast gene coding sequences comprising gene expression (SAGE) tags, useful

NORF genes with serial analysis for studying, monitoring and

expression (SAGE) tags, use ng phases of the cell cycle

1; Page 391; 419pp; English

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RESULT 111
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AAF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present
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                      The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase. S phase and G2/M; (2) a method (M2) for screening candidate
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                                                                                                                                                                                                                                                                                                            Example; Page 40;
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                                                                                                                                                                                                                                                                                                                                                                                                  ast gene coding sequences comprising NORF genes with serial gene expression (SAGE) tags, useful for studying, monitoring
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ng phases of the cell cycle
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progression comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a yeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. AAF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           expression varies as in M1, where a test substance which modifies expression of the yeast gene is a candidate antifungal drug; (3) (M3) for identifying human genes which are involved in cell cycle
                                                      invention.
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Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; nor previously assigned open reading frame; nonannotated ORF; serial analysis of gene expression; antifungal; tag; identific Saccharomyces cerevisiae linker; PCR primer; tag; identification SAGE;

WO200077214-A2

14-JUN-2000; 2000WO-US16223

16-JUN-1999; 9908-0335032

(UYJO ) UNIV JOHNS HOPKINS

Velculescu V, Vogelstein B, Kinzler K;

2001-061874/07

Yeast gene coding sequences comprising of gene expression (SAGE) tags, useful affecting phases of the cell cycle -NORF genes with serial analysis for studying, monitoring and

Example; Page 95; 419pp; English.

The present invention describes an isolated DNA molecule comprising coding sequence of a yeast gene selected from a group of 745 NORF (previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also are: method (1<u>M</u>) of using NORF to affect the cell

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RESULT 114
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ID AAF418
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Best Local
                                                                            Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
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Example; Page 308; 419pp; English.
                                                                                                                                                                                                                                             WPI; 2001-061874/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Saccharomyces cerevisiae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF;
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                                                                                                                                                                                                                                                                                                                              Vogelstein B,
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90.0%;
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The present invention describes an isolated DNA molecule comprising

Yeast gene coding sequences comprising of gene expression (SAGE) tags, useful

NORF genes with serial analysis for studying, monitoring and

expression (SAG)

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RESULT 115
AAF42385
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC antifungal drugs comprising: (a) contacting a test substance with a CC yeast cell; and (b) monitoring expression of a NORF gene whose CC expression varies as in M1, where a test substance which modifies the CC expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle CC progression comprising contacting human DNA with a probe which comprises CC at least 10 contiguous nucleotides of a NORF gene whose expression varies CC as in M1; and (4) a method (M4) for identifying a candidate drug as a CC member of a class of drugs having a characteristic effect on gene CC expression in a yeast cell comprising contacting a yeast cell with a CC candidate drug and monitoring expression in the yeast cell of at least 1 CC NORF gene whose expression is affected by the class of drugs. The NORF CC genes may be used to study, monitor and affect phases of the cell cycle. The methods may be used as markers of phases of the cell cycle. The methods may be used as markers of phases of the cell cycle and for identification of antifungal drugs. CC AAP33268 to AAP44064 represent SAGE tags used in the exemplification of the present invention. AAP33262 to AAP33267 represent linkers and PCR convention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local
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                                                                                                    Velculescu V,
                                                                                                                                                                                                               14-JUN-2000; 2000WO-US16223
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                                                                                                                                                                                                                                                                                  WO200077214-A2
                                                                                                                                                                                                                                                                                                                     Saccharomyces cerevisiae
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:9124
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                                                                                                    Vogelstein B,
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90.0%;
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Pred. No. 8
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RESULT 116
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CC addidate drug and monitoring expression in the yeast cell of phases of the cell cycle and for identification of antifungal drug.

CC addidate drug and monitoring expression in the yeast cell of phas
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Best Local (
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                                                                                                                                                                                                                                                                                                                            Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
WPI; 2001-061874/07
                                        Velculescu V, Vogelstein B,
                                                                                                                                                                                                                                                                                        Saccharomyces cerevisiae
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                                                                                                                         16-JUN-1999;
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                                                                                 (UYJO ) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                       NORF
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                                                                                 JOHNS HOPKINS
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                                                                                                                                                                                                                                                                                                                                                                                                                                   tag oligonucleotide
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Pred. No. 87;
                                        Kinzler K;
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Yeast gene coding sequences comprising of gene expression (SAGE) tags, useful affecting phases of the cell cycle -
                                                                                                                                            NORF genes with serial analysis for studying, monitoring and
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Example; Page 375; 419pp; English

cc antifungal drugs comprising: (a) contacting a test substance with a compression of a test substance with a compression of the yeast gene is a candidate antifungal drug; (3) a method (CC expression of the yeast gene is a candidate antifungal drug; (3) a method (CC (M3) for identifying human genes which are involved in cell cycle (M3) for identifying human genes which are involved in cell cycle (CC progression occupitions nucleotides of a NORF gene whose expression varies at least 10 contiguous nucleotides of a NORF gene whose expression varies (CC as in M1; and (4) a method (M4) for identifying a candidate drug as a condidate drug as a condidate drug and monitoring appression in the yeast cell with a cCC candidate drug and monitoring expression in the yeast cell with a cCC candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of drugs. The NORF comparison of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs.

CC AAF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by alleast 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate ac

Sequence 10 BP; 0 A; 4 C; 3 G; 3 T; 0 other;

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Вb
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                                     Query Match
Best Local
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             7 GCCAGGCCAA 16
                             Similarity 9; Conserv
                              Conservative
                                    42.0%;
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                                      Pred. No. 87;
                                             Score 8.4;
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                              Indels
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ABV84 RESULT 117

ABV84222 standard; cDNA; 10

12-DEC-2002 (first entry)

Human heat shock protein 40 (HSP40-1) SAGE tag

SAGE tag; serial analysis of gene expression; human; chronic hepatitis CH; liver tissue; hepatocellular carcinoma; cancer; tumour; HCC; expression pattern; differential expression; ss. Ω

Homo sapiens.

JP2002209591-A

30-JUL-2002

19-JAN-2001; 2001JP-0012328

19-JAN-2001; 2001JP-0012328

(KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN.

WPI; 2002-631294/68

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RESULT 118
ABV84641/c
ID ABV846
XX ABV846
XX ABV846
XX Human
XX SAGE t
KW CH; li
KW EXPRES
KW SS.
XX SS.
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JP2002209591-A
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                              Claim 37; Page 23; 139pp; Japanese
                                                                                            Human chronic hepatitis C tissue expression exasperating gene group comprises 100 high-ranking genes -
                                                                                                                                                                                                                                                                                                                                19-JAN-2001;
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                                                                                                                                                                                                  WPI; 2002-631294/68
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                                                                                                                                                                                                                                                               (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
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RESULT 119
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                            The invention relates to SAGE (serial analysis of gene expression) tags representing groups of genes which are differentially expressed in human chronic hepatitis C (CH) liver tissue or hepatitis C-induced hepatocellular carcinoma (HCC) compared with normal human liver tissue. The SAGE tags of this invention consist of a sequence of 10 nucleotides located downstream of the 5'-CATG-3' sequence motif lying nearest to the polyA region of cDNAs derived from a variety of genes. These tags serve
                                                                                                                                                                                                                                                                                                                                                                             Human chronic hepatitis C tissue expression exasperating gene group comprises 100 high-ranking genes -
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                                                                                                                                                                                                                                                                                                                Page
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to uniquely identify each transcript

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RESULT 120
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Best Local &
                                                                                                         The invention relates to SAGE (serial analysis of gene expression) tags representing groups of genes which are expressed in activated human Thl and/or Th2 cells. The SAGE tags of this invention consist of a sequence of 10 nucleotides located downstream of the 5'.CAG-3' sequence motif lying nearest to the polyA region of cDNAs derived from a variety of genes. These tags serve to uniquely identify each transcript and can thus be used to analyse the pattern of gene expression in particular cell types. The invention also relates to proteins encoded by the genes expressed in Th1 and/or Th2 cells, antibodies against these proteins, and inhibitors of the expression of groups of genes that are expressed in either or both the two cell types. Groups of genes expressed in Th1 and/or Th2 cell types may be used for the diagnosis and treatment of Th1 and Th2-related disorders. Sequences ABV78140-ABV78389 are SAGE tags
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 10; Page 9; 60pp; Japanese
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                                                                            are most highly expressed in
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Acid Transporter), Member 4 isogenes, for improving efficiency
eliability in drug development for treating cancers -
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                                               polymorphisms
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                                                                                                       The invention relates to an isolated polynucleotide (I) comprising a CC first nucleotide sequence which comprises solute carrier family 1 (glutamate/neutral amino acid transporter), member 4 (SLC1A4) isogenes (II) and an isolated polypeptide (III) comprising an amino acid sequence (III) and an isolated polypeptide (III) comprising an amino acid sequence (which is a polymorphic variant of a reference sequence for SLC1A4 protein. Also described are methods for: (1) haplotyping or genotyping CC SLC1A4 gene of an individual; (2) predicting an association between a trait and at least one haplotype or haplotype pair of SLC1A4 gene. (III) Is useful cin screening for drugs targeting (III) that are useful for treating cancer, neurological diseases and amino acid transporter disorders. The methods are useful for improving the efficiency and reliability of several steps in the discovery and development of drugs for treating canced by the pharmaceutical research scientist to validate SLC1A4 as a condidate target for treating a specific condition or disease predicted to be associated with SLC1A4 activity, e.g. cancer, neurological diseases cand amino acid transporter disorders, and in the design of clinical trials for treating a specific condition of disease associated with SLC1A4 activity, e.g. cancer, neurological diseases trials for treating a specific condition of disease associated with SLC1A4 activity, e.g. cancer, neurological diseases and amino acid transporter disorders, and in the design of clinical trials for treating a specific condition of disease associated with SLC1A4 antibody is useful in diagnostic, und theraneutic mathods an antibody is useful in diagnostic.
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                                  SLCIA4 gene allele-specific oligonucleotides, oligonucleotides and related PCR primers used nucleotide polymorphisms (SNP) of the gene.
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                                                                                               prognostic and therapeutic methods. ABK95761-ABK95877 represent
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The present sequence is that of an oligonucleotide used for assaying a polymorphism in the zinc finger protein homologous to zfp-36 in mouse (ZFP36) gene of the invention. The specification describes a newly cisolated polymucleotide comprising a sequence which is a polymorphic curis fragment and its encoded protein. The ZFP36 gene (see ABA97001) or its fragment and its encoded protein. The ZFP36 polymorphic curis fragment and its encoded protein. The ZFP36 polymorphic curis fragment and its encoded protein. The ZFP36 polymorphic curis fragment and its encoded protein. The ZFP36 polymorphic curis fragment and its encoded protein. The ZFP36 polymorphic curis fragment and its encoded protein. The ZFP36 polymorphic curis fragment in the ZFP36 polymorphic curis for the ZFP36 polymorphic curis for screening drugs targeting the ZFP36 polymorphic curis for the discovery and development of drugs for treating diseases associated with ZFP36 activity, e.g., curis for an individual gives useful information for validating ZFP36 as a curis for an individual gives useful information for validating ZFP36 as a curis for an individual gives useful information predicted to be candidate target for treating a specific condition predicted to be candidate target for treating a specific condition predicted to be candidate target for information used for developing diagnostic tests and cherapeutic methods. A recombinant non-human curis specific formats and therapeutic methods. A recombinant non-human curis formation in vivo drug screening and testing. Antipore in the target for specific and primers in the target region and primers.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel polymorphic variants of zinc finger protein homologous to zfp-36 in mouse gene, useful in studying expression and function of the protein, useful for screening drugs to treat diseases e.g. rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antiarthritic; drug sciening, account; haplotyping; genotyping; autoimmune disease; rheumatoid arthritis; haplotyping; genotyping; allele specific oligonucleotide, ASO; single nucleotide polymorphism; allele specific oligonucleotide, ASO; single nucleotide polymorphism; allele specific oligonucleotide, ASO; single nucleotide polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 14; 60pp;
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                                                                                                                                                                                     The invention relates to single nucleotide polymorphisms in the gene cancoding the human B-factor properdin protein (BF). A method for haplotyping the BF gene in an individual comprises identifying the nucleotide at one or more polymorphic sites and determining whether one of the copies of the gene is defined by one of the BF haplotype given in the specification or whether both copies are defined by a haplotype pair. This method is useful in genotyping, whereby all possible haplotype pairs can be assigned to specific genotypes. An association between a trait and a haplotype or haplotype pair of the BF gene can be identified by comparing the frequency of the haplotype pair in a population exhibiting the trait with the frequency of the haplotype pair in a population and indicates the trait is associated with the haplotype pair in a reference population, where a higher haplotype or haplotype pair in a reference population, where a higher haplotype for studying the expression and function of BF, for use in screening for candidate drugs to treat diseases related to BF activity, such as diabetes and systemic lupus erythematosus. Sequences ABK6450-ABK64105 represent allee-specific PCR primers used to detect human BF gene
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                                                                                                                Sequence
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Matches 9
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                                                              ABK23567;
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                                                                                                                       ABK23567 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        relates to novel single nucleotide polymorphisms an GTPase activating protein 1 (RANGAP1) gene
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Pred. No. 87;
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09-APR-2002

(first entry)

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                                                                                                                     Human; retinaldehyde-binding protein 1; ss; RLBP1; haplotype; primer; genotyping; probe; autosomal recessive retinitis pigmentosa; arRP; PC chromosome 15q26; transgenic; ASO; allele specific oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A new nucleic acid library of myc-dependent downstream genes capable of supporting a neoplastic characteristic of cancer is useful to find new therapies and diagnoses for cancer -
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29-JUN-2000; 2000EP-0202284.
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29-MAY-2001; 2001WO-US17252.
                               06-DEC-2001
                                                                                                                                                                                Retinaldehyde-binding protein 1 ASO primer extension
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                                                                                                                                                                                                                                               ABK24232;
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                                                                                        Homo sapiens
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                                                                                                                                                                                                              entry)
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90.0%;
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protein 1 gene, useful for studying RLBP1 protein for use in screening to RLBP1 activity -
                                                                                                                                                                                                                                                                         New genetic variants having polymorphisms in the retinaldehyde-binding protein 1 gene, useful for studying the function of and for expressing RLBP1 protein for use in screening drugs for treating diseases related
                                                                                                                                                                                                                                                         Claim 18; Page 14; 107pp; English
                                                                                                                                                                                                                                                                                                                                   26-MAY-2000; 2000US-207618P.
                                                                                                                                                                                                                                                                                                                       GENAISSANCE
                                                                                                                                                                                                                                                                                                           Kazemi A,
                                                                                                                                                                                                                                                                                                                       PHARM INC
                                                                                                                                                                                                                                                                                                            Koshy B;
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CC genes and haplotypes of the retinaldehyde-binding protein I (RIBPI) gene are referred to as P81\_24 to designate the order in which they are clocated in the gene. Also included are methods for haplotype pair for the RIBPI gene of an individual, a method for predicting a dentifying an association between a trait and at least one haplotype pair for the RIBPI gene of an individual, a method for predicting a cleantifying an association between a trait and at least one haplotype pair of the RIBPI gene of an individual, a method for predicting a palotype pair of the RIBPI gene of an individual, a method for predicting a palotype pair of the RIBPI gene of an individual, a method for predicting of palotype pair of the RIBPI gene of an individual, a method for predicting of palotype pair of the RIBPI gene of an individual comprising a set of oligonucleotides designed to genotype each of palotype pair of the RIBPI gene of an individual comprising a set of oligonucleotides designed to genotype each of palotype produced by the polymorphic variant sequence or expresses an RIBPI protein encoded by the polymorphic variant sequence or expresses a RIBPI protein encoded by the polymorphic variant sequence or expresses a RIBPI protein encoded by the polymorphic variant sequence or expresses an RIBPI protein for genetype gene that is a polymorphic comprising an amino acid sequence that is a polymorphic curiant sequence, an isolated complement process in control of RIBPI and in studying the expression and function of RIBPI gene is useful in studying the expression and function of RIBPI gene is custoff and in expression and function of RIBPI, and in expression and function of RIBPI and in expression and function of RIBPI and in the gene of a sequence is a sequence for compounds cargeting RIBPI control of sease.

CC useful in improving the expression and function of sease are useful for designing clinical trials of candidate drugs for sceening completed to RIBPI and trials of candidate drugs for sceening completed to a polymorphic a polymorphic RLBP sequence, using invention relates to an isolated polynucleotide, which comprises the primer extension method 1 (RLBP1) ge...

Sequence 10 BP; 3 A; 4 C; 3 G; 0 U; 0 other;

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90.0%;
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                             Mismatches
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RESULT 128 AAS99275 ID AAS991

AAS99275 standard; 10

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RESULT 129
AAS16750/c
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                                                                                                                                                                                                                                            cc nucleotide at one or more polymorphic sites and determining whether one cof the copies of the gene is defined by one of the F12 haplotypes given in the specification or whether both copies are defined by a haplotype con in the specification or whether both copies are defined by a haplotype pair. This method is useful in genotyping, whereby all possible haplotype pairs can be assigned to specific genotypes. An association between a comparing the frequency of the haplotype pair of the F12 gene can be identified by comparing the frequency of the haplotype or haplotype or haplotype or haplotype or comparing the frequency of the haplotype or haplotype or comparing the frequency of the haplotype or compulation indicates the trait is associated with the haplotype pair in a reference population, where a higher haplotype or computation indicates the trait is associated with the haplotype or haplotype pair. F12 and its corresponding DNA are used corresponding to treat disorders related to F12, for use in screening for comparing the expression and function of F12, for use in screening for comparing artery disease, liver disease, spontaneous abortion, Alzheimer's corresponded to the first of the sequences are also useful for studying the effect of variation on the biological activity of F12 as well as on the binding affinity of candidate drugs targeting F12. Sequences AAS99209-AAS99305 represent problems, primers and primer tails used to detect F12 gene polymorphisms.
                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to single nucleotide polymorphisms in the gene encoding the human coagulation factor XII (F12) polypeptide. A metho haplotyping the F12 gene in an individual comprises identifying the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel isolated human coagulation factor XII polynucleotide, F12 use for treatment of e.g. coronary artery disease, comprises a sequence which is a polymorphic variant of a reference sequence for F12 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             single nucleotide polymorphism; genotyping; gene therapy; drug screening; coronary artery disease; liver disease; spontaneous abortion; cardiant; Alzheimer's disease; blood coagulation; hepatotropic; neuroprotective;
                                                                                                                                                                                                                                          Sequence 10
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RESULT 130
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                                               Human maturation/activation dendritic cell expression gene tag #45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New haplotypes of the human apolipoprotein A-IV gene, useful to diagnose and treat disorders associated with its abnormal expression function such as coronary artery disease -
                                                                                                              ABL42671;
                                                                                                                                              ABL42671 standard; cDNA; 10
                                                                                                                                                                                                                                                                                                                                                             Sequence 10 BP; 0 A; 5 C; 3
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atherosclerosis; PCR primer; primer extension.
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                                                                                                                                                                                                                                                                                             Conservative
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90.0%;
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                                                                                                                                                                                                                                                                                           0; Mismatches
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Pred. No. 8
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Human; maturation/activation dendritic cell maturation; activation; dendritic cell; ss.

expression gene; tag;

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RESULT 131
ABV62380
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Best Local
                                                                                                                                                                                                                                      Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      highest expression among the genes expressed in human maturation/ activation DC. Also described are: (1) a protein expressed by the above human maturation/activation DC expression gene; (2) an antibody against the protein; and (3) an antagonist against the expression of each gene belonging to the above gene group. The gene group is useful for the treatment and the diagnosis of various human diseases related to human DC. ABL42627 to ABL42926 represent specifically claimed human
In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                               03-JAN-2001; 2001DE-1000127
                                                                                                                                       20-DEC-2001;
                                                                                                                                                                  11-JUL-2002
                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                           Human skin
                                                                                                                                                                                                                                                                                                                    21-OCT-2002
                                                                                                                                                                                                                                                                                                                                             ABV62380;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 10 BP; 1 A; 3 C; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         maturation/activation DC expression gene tags from the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim
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                                                                                                                                                                                         WO200253774-A2
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                                                                                        (HENK ) HENKEL
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                                       2002-590638/63
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                                                                Conradt M,
                                                                                        KGAA
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                                                                Hofmann
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DT 21-
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Best Local (
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                                              The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or
                                                                                                                                                                                                         e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                         Claim
                                                                                                                                                                                                                        In vitro identification of skin-expressed homeostasis and identifying cosmetic or ph
                                                                                                                                                                                                                                                                                                                                                                                   03-JAN-2001; 2001DE-1000127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABV69801;
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                                                                                                                                                                                                                                                                                                                                                  (HENK ) HENKEL KGAA
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9; Conserv
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90.0%;
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                                                                                                                                                                                                                                                                                                                Hofmann
                                                                                                                                                                           German.
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Pred. No. 1.1e+02;
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                                                                                                                                                                                                                          ed genes, useful for pharmaceutical agent
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promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; sc

atopic dermatitis; acne; seborrhea; lupus erythematosus;

scleroderma;

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RESULT 133
AAX54916
ID AAX549
XX AAX549
XX AAX549
XX AAX549
XX AAT18e
AC C/EBP-
XX Ant18e
KW Ant18e
KW impair
KW colon
KW colon
KW colon
KW colon
KW colon
KW colon
KW pulmon
KW colon
KW posta
OS Synthe
YN WO9913
XX Synthe
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XX O9-JUN
PR 17-SEP
XX O9-JUN
PR 09-JUN
PR 0
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Best Local S
Matches
                                                                                                              The specification describes antisense oligonucleotides (AAX52869-X55271) directed against at least 2 mRNAs selected from target genes, coding and non-coding regions of RNAs corresponding to target genes, gene initiation codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'-end and the juxta-section between coding and non-coding regions and all segments of RNAs encoding proteins associated with one or more diseases, conditions or mixtures. The antisense oligonucleotides may be derived from sequences AAX55272-74. These multiple target oligonucleotides (specifically AAX55180-271) can be used for the antisense treatment of diseases and conditions. Typical diseases and conditions are those associated with impaired respiration and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense oligonucleotide; multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vasoconstriction; inflammation; allergic rhinitis; acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary hypertension; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer; colon cancer; breast cancer; lung cancer; pancreatic cancer;
                       inflammation, including lung diseases, pulmonary vasoconstriction, inflammation, allergic rhinitis, acute asthma, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 68; 120pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense oligonucleotides used in treatment of, e.g. pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-JUN-1998;
17-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         prostate cancer; ss.
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hypertension,
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97US-0059160.
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Pred. No. 1.1e+02;
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                                                                                                                                                                                              The present invention describes low adenosine (A) content antisense oligonuclectides and compositions (I) comprising them. In the antisense oligonucleotides the A is replaced by a 'Universal' or alternative base. (I) can have respiratory, bronchodilator, antiinflammatory, analyssic, immunosuppressive, antiasthmatic, hypotensive and cytostatic activities.
                                                                                                                                                                                                                                                                                                                                                                                                                            Low adenosine (A) content antisense oligonucleotides which trigger adenosine receptors during metabolism, useful e.g. cancers and respiratory obstructions -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (NYCE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              respiratory obstruction, pulmonary obstruction; impeded respiration; surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pain; cystic fibrosis; allergic rhinitis; pulmonary hypertension; emphysema; pulmonary transplantation rejection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Low adenosine antisense oligonucleotide; phosphorothicate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 lymphomas, carcinomas e.g. colon cancer, breast cancer, lung canc pancreatic cancer, hepatocellular carcinoma, kidney cancer, melar hepatic metastases, as well as all types of cancers which may met or have metastasized to the lungs, including breast and prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-APR-1999;
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Pred. No. 1.1e+02;
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immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. The antisense oligonuclectides and (I) can be used to down-regulate the expression and or activity of target polypeptides associated with lung/respiratory disorders and malignancies, such as stimulating and activating peptide factors and transmitters, transcription factors, immunoglobulins and antibodies, antibody receptors, cytokines and chemokines, endogenously produced specific and non-specific enzymes, immunoglobulins and antibodies antibodies antibodies and non-specific enzymes.

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The present invention describes a new composition antisense oligonucleotide (ON) with low adenosine
                                                                                                                                                                           New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstriction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   condition selected from pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome (RDS), pain, cystic fibrosis (CP), allergic rhinitis (AR), pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD), pulmonary transplantation rejection, pulmonary infections, bronchitis, and/or cancer. AAF18434 to AAF21543 represent human polynucleotide fragments and antisense oligonucleotides used in the exemplification of the control of 
                                                                                        Disclosure; Page 523; 1343pp; English
                                                                                                                                                                                                                                                                                                          WPI; 2000-205971/18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-AUG-1998;
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Pred. No. 1.1e+02;
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comprising an (up to 15%), which
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                release of deoxyadenosine which activates adenosine receptors causing bronchoconstriction and inflammation. AAA32313 to AAA35312 represent the nucleotide sequences given in the sequence listing from the present invention, which correspond to SEQ ID NO:1 to 2815, and then the last 185 sequences are also called SEQ ID NO:1 to 185, but the sequences differ from the previously named sequences. SEQ ID NO:11 to 1680 (AAA32323 to AAA33992) are specifically claimed ONs from the present invention. N.B. Sequences given in the disclosure of the present invention do not match up with their corresponding SEQ ID NO: sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  asthma, impeded respiration, respiratory distress syndrome, pain, cystifibrosis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukaemias, lymphomas, carcinomas, and cancers which may metastasise to the lungs, including breast and prostate cancer. The reduction of the adenosine content of the ONs reduces side effects. The A-containing ONs break down with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     targets nucleic acids involved in bronchoconstriction, allergies, and/or inflammation. The ON can have antiinflammatory, antiallergic, antiasthmatic, cytostatic and analgesic activities. The compositions are useful for the treatment of diseases associated with inflammation, impaired airways, including lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating e.g. ischaemic conditions, pulmonary vasoconstriction, allergies,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUP-TF binding site; inverted repeat; minimal packaging element; adenovirus; chicken ovalbumin upstream promoter transcription factor; repressor binding site; DNA delivery; recombinant virus; ds
                                                                                           Regulating adenoviral packa
sites that allow selective
                                                                                                                                                            WPI; 2000-052657/04.
                                                                                                                                                                                                                                                                               15-APR-1998;
05-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUP-TF inverted repeat binding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAZ59919 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 11 BP; 0 A; 4 C; 6 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               given in the sequence listing.
                                                                                                                                                                                                                                          (UYNY ) UNIV NEW YORK STATE RES
                                                                                                                                                                                                                                                                                                                                            15-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4 CGGGCCAGGC
                                                                                                                                                                                                  ָס,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity 9; Conserv
                                                                                                                                                                                                    Schmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                               98US-0081867
98US-0088321
                                                                                                                                                                                                                                                                                                                                            99WO-US08294
                                                                                                                                                                                                    SI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             42.0%;
                                                                                           packaging by inco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11
                                                                                                                                                                                                  Ostapchuk PH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 8.4; DB 1;
Pred. No. 1.1e+02;
                                                                                                by incorporation of repressor binding ression of packaging, used for gene
                                                                                                                                                                                                                                            FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      site.
                                                                                                                                                                                                    Erturk
                                                                                                                                                                                                    m
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                              for gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    pain, cystic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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The invention relates to the regulation of adenoviral packaging

Example 5;

Page 39; 71pp; English

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local (
                                     Claim 12; Page 8;
                                                                                            Determining presence of a tumor cell or angiogenesis, effectiveness of treatment, by detecting the presence
                                                                                                                                                     WPI; 2002-668396/72.
                                                                                                                                                                                                                                                                  23-JAN-2001; 2001EP-0200228.
28-SEP-2001; 2001EP-0203703.
28-SEP-2001; 2001US-325722P.
                                                                                                                                                                                         Van Der Kuyl AC,
                                                                                                                                                                                                                                                                                                                                             23-JAN-2002; 2002EP-0075264
                                                                                                                                                                                                                                                                                                                                                                                  24-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                        EP1225233-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; Kaposi's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kaposi's Sarcoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABQ81854 standard; DNA; 11 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 11
                                                                                                                                                                                                                              (AMST-) AMSTERDAM SUPPORT DIAGNOSTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chod of the invention comprises propagating an adenoviral containing a repressor binding site, in the absence of t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          9;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     packaging.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                          detect and monitor treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sarcoma; tumour; angiogenesis; tag;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SAGE library Tag No.4 SEQ ID
                                     38pp;
                                                                                                                                                                                         Cornelissen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              42.0%;
                                   English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 8.4; DB 1;
Pred. No. 1.1e+02;
                                                                                        detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                              B۷
                                                                        t of
                                                                        presence
Kaposi's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 88
                                                                        of marker
Sarcoma -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
                                                                                          genes
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has a tumour cell or site of angiogenesis, or if a treatment is effective in changing angiogenesis or changing a status of a set of target cells, comprising determining if a sample of the subject has an expression product of at least one marker gene. Also described is a compound capable of altering the expression or activity of Keratin 14, TIE 1, Salioadhesin or Siglec in a cell. Peripheral blood mononuclear cell (PBMC) expressed Keratin 14, TIE 1, Salioadhesin or Siglec, and kits containing them from the present invention can be used in a diagnostic method, particularly as an indicator of angiogenesis or to determine presence of a tumour cell. The method of the invention is suitable to determine within a few days if a certain treatment against Kaposi's Sarcoma is successful. Ap081851 to
                                                          present invention.
                                                                               ABQ82006 represent nucleotide sequence used in
                                                                                     the exemplification
                                                                                        of the
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Sequence 11 BP; 3 A; 4 C; 4 G; 0 U; 0 other;

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            S
                                  Query Match
Best Local
                              Matches
              11
                            Similarity 9; Conserv
              GGCCAAAGTC
GGCCAAAGGC
                             Conservative
10
              20
                                   42.0%;
                            0
                                   Score 8.4; DB Pred. No. 1.1e+
                              Mismatches
                                   .1e+02;
                                         Length 11;
                             Indels
                             0
                             0
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RESULT 138
ABV62551/c
ID ABV625
XX ABV625
XX ABV625
XX Human
DE Human
XX Human;
KW immuno
KW psoria
XX Jenes
PN W02002
XX Jenes
PN W02002
XX HENK
XX Peters
XX Peters
XX In vit
PT homeos
PT In vit
PT homeos
PT In vit
PT e-g. s
XX Discle

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
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11-JUL-2002 WO200253774-A2

03-JAN-2001; 2001DE-1000127 20-DEC-2001; 2001WO-EP15179

In vitro identification of skin-expressed homeostasis and identifying cosmetic or pl (HENK ) HENKEL 2002-590638/63 KGAA 3 Hofmann ed genes, useful for de pharmaceutical agents

determining s against

cancer

The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of skin. The present sequence is that of a human expressed sequence tag Page 35; 1345pp; German. scleroderma; the

The present invention describes a method

for

determining

if an individual

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RESULT 139
ABV62704
ID ABV627
XX ABV627
XX ABV627
XX ABV627
XX ABV627
XX Human;
KW Human;
KW Immuno
KW Jesoria
XX Homo s
XX Homo s
XX Homo s
XX Homo s
XX WO2002
XX In JUL
PF 20-DEC
XX WPI; 20-DEC
XX WP
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                                                                                                      Query Match
Best Local S
Matches 9
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Best Local
                                                                                                                                                                                                                                                    in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma expressed sequence tag (EST) of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
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                                                                                                                                                                                                             Sequence 11 BP; 1 A; 5 C; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 39; 1345pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Petersohn D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-JAN-2001; 2001DE-1000127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HENK ) HENKEL KGAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10
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                                                                                                      Similarity 9; Conserv
GGCCCAGGCC 10
                                               GGGCCAGGCC 14
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                                                                                                         Conservative
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                                                                                                                               42.0%;
90.0%;
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90.0%;
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                                                                                                                                                                                                             G; 1 T; 0 other;
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                                                                                                                             Score 8.4; DB 1;
Pred. No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 8.4; DB 1;
Pred. No. 1.1e+02;
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                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            <u>..</u>
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                                                                                                                                                         Length 11;
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RESULT 140

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RESULT 141
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               Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
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                                                                                 Human skin EST
                                                                                                                    21-OCT-2002
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Pred. No. 1.1e+02;
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     Petersohn D,
                                                          (HENK ) HENKEL KGAA
                                                                                                               03-JAN-2001; 2001DE-1000127.
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ying cosmetic or pharmaceutical agents
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Pred. No. 1.1e+02;
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neurodermatitis;
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The invention relates to in vitro identification in the skin of humans or animals by subjecting a encoded factors from skin, to serial analysis of

coded factors from skin, to serial ar as to identify skin-expressed genes

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cation (M1) of genes expresse tring a mixture of genetically rsis of gene expression (SAGE) quantify their expression.

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RESULT 144
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                              The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
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                                                                                                                                                                                                                                                       Disclosure; Page 129; 1345pp; German.
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RESULT 146 ABV66964/c

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Matches
                                                                                  Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
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                             WO200253774-A2
                                                          Homo sapiens.
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                                                                                                                                                                                                                                                          Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
In vitro identification of skin-expressed genes, useful for de homeostasis and identifying cosmetic or pharmaceutical agents
                                          WPI; 2002-590638/63.
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Pred. No. 1.1e+02;
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.g. skin cancer

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RESULT 149
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EN WO2002
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                                                                                   The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or
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                         promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; sc
                                                                                                                                                                                                                                                                                                               Disclosure;
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Pred. No. 1.1e+02;
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Score 8.4; DB 1; Pred. No. 1.1e+02;

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XX Human;
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                                                     The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosaces; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention.
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(EST)
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ed genes, useful for determining pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .1e+02;
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RESULT 152
ABV99138/c
ID ABV691
XX
AC ABV691
XX
DT 21-OCT
XX
DE Human
XX
KW Human;
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                                                                                                                                                                                                                                                                                                      RESULT 151
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Best Local (
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                                                                                                                                                                                                                                                                                                                       in the skin of humans or animals by subjecting a mixture of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis and to test agent (A) that maintains or disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence term (EST) of the invention.
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Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic;
                             Human skin
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                                                          21-OCT-2002
                                                                                      ABV69138;
                                                                                                                 ABV69138 standard; cDNA;
                                                                                                                                                                                                                                                                                                      Sequence 11 BP; 3 A; 4 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 215; 1345pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Petersohn D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-JAN-2001; 2001DE-1000127
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                                                                                                                                                                                                                                              Similarity 9; Conserv
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                             EST 6924
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                                                        (first entry)
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90.0%;
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Pred. No. 1.1e+02;
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RESULT 153
ABV69411/c
ID ABV694
XX ABV694
XX 21-OCT
XX Human
XX Human;
KW Human;
KW psoria
XX Psoria
XX DSO Homo s
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                                                                                                                                                                                                                                                                                                                             Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                               03-JAN-2001; 2001DE-1000127
                                                                                       20-DEC-2001; 2001WO-EP15179
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human skin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          e.g. skin cancer
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                                                                                                                                                                                                                                                                        Homo sapiens
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90.0%;
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Pred. No. 1.1e+02
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                                                                                                                                                                                    Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                       WO200253774-A2
                                                                                                                                                                                                                    Human skin
                                                                                                                                                                                                                                                  ABV69528
                                                                                                                                                                                                                                                                  ABV69528 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 226; 1345pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             homeostasis and identifying
             Disclosure; Page 229; 1345pp; German
                                   In vitro identification of skin-expressed homeostasis and identifying cosmetic or pl
                                                            WPI; 2002-590638/63
                                                                                                         03-JAN-2001; 2001DE-1000127
                                                                                                                        20-DEC-2001; 2001WO-EP15179
                                                                                                                                                                      Homo sapiens.
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                                                                                          (HENK ) HENKEL
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                              cancer
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                                                                           Conradt
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Pred. No. 1.1e+02
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                                    pharmaceutical agents
                                           genes, useful for determining
                                                                                                                                                                                                                                                                                                                                                   Length 11;
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The invention relates to

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genes expressed

Sequence 11

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RESULT 155
ABV69972/c
ID ABV699
XX ABV699
XX ABV699
XX ABV699
XX ABV699
XX INTERPORT 21-OCT
XX Human;
KW Immuno
XX Human;
KW Immuno
XX Homo s
XX WO2002
XX Homo s
XX WPI;
PF 20-DEC
XX WPI;
PF 20-DEC
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Best Local S
Matches 9
                                                        The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 24; Page 247; 1345pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
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suppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis;
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Pred. No. 1
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.1e+02;
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RESULT 157
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ABV70125
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Matches
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Best Local :
                                                                                                                                                                                                    The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; chthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention.
  ABV70458 standard;
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                                                                                                                                                                           Sequence 11 BP; 1 A; 5 C; 4
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                                                                                                                                                                                                                                                                                                                                                                                                                         In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Petersohn D,
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                                                                                                                                                                                                                                                                                                                                                                                    Page 252;
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                                                                                                                        Conservative
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90.0%;
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Pred. No. 1.1e+02;
0; Mismatches 1
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.1e+02;
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ABQ86459
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Best Local S
Matches 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; acopic dermatitis; acne; seborrhea; lupus erythematosus; rosacca; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention.
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                                              Human
                                                                                          10-SEP-2002
                                                                                                                                     ABQ86459
                                                                                                                                                                               ABQ86459 standard;
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                                         skin stress/ageing related EST
  skin
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ageing;
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  skin
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  stress;
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                                                                                                                                                                                 ВP
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                                                                                                                                                                                                                                                                                                                                                                                                      Score 8.4; DB 1;
Pred. No. 1.1e+02;
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  EST;
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  expressed
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  sequence
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RESULT 159
ABQ86573/c
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Best Local S
Matches 9
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         gene
                                                                                                                                      03-JAN-2001; 2001DE-1000121
                                                                                                                                                                                                                                                                  Human; skin ageing; skin stress; EST; expressed sequence tag;
                                                                                                                                                                                                                                                                                            Human skin stress/ageing related
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                     Identifying genes involved in skin stress and screening for cosmetic or therapeutic agents,
                                                                                    Petersohn D,
                                                                                                                                                               20-DEC-2001; 2001WO-EP15178.
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                                                                                                                                                                                                                  WO200253773-A2
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                                                                                                               (HENK ) HENKEL KGAA.
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          expression
                                                            2002-528865/56
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90.0%;
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Pred. No. 1.1e+02;
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based on differenti
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RESULT 160
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                           expression. (A) comprises protein or mRNAs or their fragments. (M1) is useful for: identifying markers of skin ageing and/or stress; determining skin ageing and/or stress; and identifying or determining the effects of pharmaceutical or cosmetic agents for control of skin ageing. The present sequence is one of a group of human skin ageing/stress related expressed sequence tags (ABQ86246-ABQ87680) of the invention.
                                                                                                            The invention relates to identifying (M1) genes in vitro that, in humans or animals, are important for skin ageing and/or skin stress by serial analysis of gene expression between mixtures of transcribed and optionally translated, genetically encoded factors (A) obtained from young and aged skin, to identify that genes that show strong differential expression. (A) comprises protein or mRNAs or their fragments. (M1) is
                                                                                                                                                                                                                                Claim 8; Page 75; 325pp; German.
                                                                                                                                                                                                                                                                              Identifying genes involved in skin stress and ageing, useful e.g. screening for cosmetic or therapeutic agents, based on differentia
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                                                                                                                                                                                                                                                                                                                                                                 Conradt M,
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Pred. No. 1.1e+02;
0; Mismatches 1
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Sequence 11

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    RESULT 162
ABL92021
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Best Local S
Matches 9
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Best Local S
Matches 9
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    30-MAY-2002
                                            ABL92021;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Petersohn D,
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Pred. No. 1.1e+02;
0; Mismatches 1.
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Pred. No. 1.1e+02;
D; Mismatches 1
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tumour angiogenesis;

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Query Match
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Matches 9
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Human; endothelial cell; EC; tumour endothelial cell; Tumour endothelial marker; normal endothelial marker; pan-endothelial marker; polycystic kidney disease; pso diabetic retinopathy; rheumatoid arthritis; tumour ang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an isolated molecule comprising an antibody variable region which specifically binds to an extracellular domain c tumour endothelial marker (TEM) protein selected from ABB90732, ABB90 ABB90749, ABB90750 and ABB90769. The antibodies which bind to TEM
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11-AUG-2000; 2000US-224360P
11-APR-2001; 2001US-282850P
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polycystic kidney disease; diabetes; retinopathy; rheumatoid arthritis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>,,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 8.4; DB 1;
Pred. No. 1.1e+02;
0; Mismatches 1
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                                             psoriasis;
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                                                                                       PEM;
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AAA80792
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Best Local Similarity
Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a novel method for the isolation of endothelial cells (ECs), and the identification of genes expressed in normal and tumour ECs. Tumour endothelial marker (TEM), normal endothelial marker (NEM), and pan-endothelial marker (PEM) genes are identified in human ECs. The human EC marker proteins and the polynucleotide sequences encoding them are useful for detecting, diagnosing or treating tumours as well as polycystic kidney disease, diabetic retinopathy, rheumatoid arthritis, and psoriasis. They are also useful for inhibiting neoangiogenesis or tumour angiogenesis, for inducing an immune response to tumour endothelial cells in a patient, or for identifying candidate drugs for treating tumours. ABX71828-ABX71999 represent DNA tags for human PEM, TEM or NEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New purified human transmembrane protein, designated as tumour endothelial marker (TEM) 3, useful for detecting, diagnosing or treating tumours, polycystic kidney disease, diabetic retinopathy,
                                                                                                                                                                  A. thaliana primer walking octamer SEQ ID NO:
                                                                                                                                                                                                                                                     AAA80792 standard; DNA; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 11 BP; 4 A; 3 C; 3 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-APR-2001; 2001US-282850P.
06-FEB-2002; 2002US-354262P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       neoangiogenesis; immune response; cytostatic; antidiabetic;
ophthalmological; antirheumatic; antiarthritic; antipsoriatic;
 15-APR-1996;
                          21-MAY-1997;
                                                       04-JUL-2000.
                                                                                   US6083695-A
                                                                                                          Arabidopsis thaliana
                                                                                                                                      Primer walking;
                                                                                                                                                                                               24-NOV-2000
                                                                                                                                                                                                                         AAA80792;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            rheumatoid arthritis or psoriasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-093016/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Carson-Walter E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-APR-2002; 2002WO-US08253
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                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                               (first entry)
96US-0632782
                          97US-0859954.
                                                                                                                                       octamer; primer;
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                                                                                                                                                                                                                                                                                                                                                                                             42.0%;
90.0%;
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Pred. No. 1.1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                       DNA sequencing; PCR;
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ARESULT 165
AAV63091/c
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AC Tof; b
KW Tof; b
KW Cancer
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AXX Concer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                  Barker N,
Sparks AB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention describes a novel method for sequencing an unknown DNA molecule which comprises selecting a library primer from an octamer oligonucleotide library consisting of 48 8-bp sequences and corresponding complementary sequences, where the library primer is complementary to a known sequence adjacent to the unknown sequence or is complementary to a sequence in a known extension product. The method is useful for DNA nucleotide sequencing, in PCR, and in other processes which make use of primers. The octamers are used to identify coding sequences. Primer walking using the octamer libraries is advantageous over other sequencing methods because it does not require multiple cloning steps nor subsequent template preparations, and it is a directed and methodical approach. AAA80688-A81253 represent the octamer primers used in the primer walking method of the invention.
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Intron-free DNA encoding Tcf-4 protein - useful identifying drugs for treating FAP patients, or increased risk of developing cancer
                                                                                                                                                                                                                                                                                                                                     20-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                         20-MAR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tcf; beta-catenin; human; drug; familial adenomatous polyposis; FAP; cancer; adenomatous polyposis coli; APC; neoplastic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human Tcf mutant DNA motif
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                                                                                                            WPI; 1998-531569/45.
                                                                                                                                                                                                                                               (UYUT-) RIJKSUNIV UTRECHT. (UYJO) UNIV JOHNS HOPKINS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12 GCCAAAGT 19
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8; Conserv
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                                                                                                                                                             Clevers H,
Vogelstein
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Pred. No.
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o. 5.8e+02;
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                            for, e.g.
patients with
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cc with beta-catenin in a method of identifying candidate drugs for used coin familial adenomatous polyposis (RAP) patients, or patients with cc increased risk of developing cancer. The protein can also be used to cetermine the presence or absence in a cell of wild type adenomatous coin (APC) gene or a downstream protein in the APC cranscription regulatory pathway. This method involves introducing a ccf-responsive reporter gene into the cell, and measuring transcription of the reporter gene where a cell which supports active transcription of the APC transcription regulatory pathway. The protein can also be used to comprise the cell, and measuring transcription of the APC transcription regulatory pathway. The protein can also be used to a method of diagnosing cancer in a sample suspected of being cancer associated with FAP, comprising administering to the patient a colypeptide comprising a portion of the APC sequence, comprising the collection of the APC cancer associated with FAP, comprising administering to the patient and collection of the APC sequence, comprising the
   RESULT 166
AAX99683/c
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Best Local S
Matches 8
The invention relates to mammalian transcription factors Tcf-3 or Tcf-4 protein or their derivatives. The Tcf proteins of the invention comprise a beta-catenin binding domain and are capable of complexing with beta-catenin in response to signaling through the Wnt/Wingless cascade. The Tcf-3 and Tcf-4 genes, proteins and antibodies against the proteins may be used to screen for candidate anticancer drugs by identifying compounds which interact with beta-catenin/Tcf-3 or beta-catenin/Tcf-4 complexes
                                                                                                                                  Disclosure; Page 4;
                                                                                                                                                                                                                                                                                                                                                   01-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                 myeloblastoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               binding
                                                                                                                                                                                                                                                                                 27-FEB-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                transactivation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mammalian;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tcf optimal response
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           an; transcription factor; Tcf-3; Tcf-4; beta-catenin; mouse;
domain; Wnt signalling; Wnt/Wingless cascade; anticancer drug;
tivation; fetal cancer; testis cancer; ovary teratoma; mammary;
                                                                                                                                                                 se transcription factors, useful for diagnosing and and screening for anticancer drugs
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8; Conserv
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                                                                  The present invention describes a zinc finger protein (I) that binds to CC a target site, comprising a first (FI), a second (F2), and a third (F3) cC zinc finger, ordered FI, F2, F3 from N-terminus to C-terminus, where the CC zinc finger sordered FI, F2, F3 from N-terminus to C2-terminus, where the CC cand a third (S3) target subsite. Also described are: (1) a second (S2), and a third (S3) target subsite. Also described are: (1) a polypeptide (CC (II) comprising (I); (2) a polynuclectide (III) encoding (I) or (II); and CC (3) designing (M) (I) involves selecting the F1 zinc finger such that the binds to the S1 target subsite, selecting the F2 zinc finger such that it binds to the S2 target subsite, and selecting the F3 zinc finger such that it binds to the S3 target subsite, thus designing (I) that binds to a target sibsite, and selecting the F3 zinc constitution of the SC target subsites and plant engineering. (I) (II) or (III) is useful in the CC subsite. (I) is useful in studying gene function, and for human the therapeutic methods to modulate the expression of a target region within
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New zinc finger protein that binds to target site, useful in studying gene function and for human therapeutics and plant engineering, comprises first, second and third zinc fingers, ordered from N- to
                                                a subject,
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1;
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in diagnostic methods for sequence specific detection of leic acid in a sample, and in assays to determined the and function of gene expression. (I) has improved affinity
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Pred. No.
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b. 5.2e+02;
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RESULT 168
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The present invention describes a zinc finger protein (I) that binds to CC a target site, comprising a first (FI), a second (F2), and a third (F3) CC zinc finger, ordered FI, F2, F3 from N-terminus to C-terminus, where the CC target site comprises, in 3-5' direction, a first (SI), a second (S2), CC and a third (S3) target subsite. Also described are: (I) a polypeptide (II) comprising (I); (2) a polyprucleotide (III) encoding (I) or (II); and CC (II) comprising (I); (2) a polyprucleotide (III) encoding (I) or (II); and CC (I) designing (M).(I) involves selecting the F1 zinc finger such that CC it binds to the S1 target subsite, selecting the F2 zinc finger such that it binds to the S2 target subsite, and selecting the F3 zinc (CC that binds to the S2 target subsite, and selecting the F3 zinc (CC that binds to the S1 target subsite, thus designing (I) CC that binds to a target site. (I) is useful for recognition of triplet (CC target subsites having the nucleotide G in the 5'-most position of the CC subsite. (I) is useful in studying gene function, and for human (CC therapeutic methods to modulate the expression of a target region within CC a subject, in diagnostic methods for sequence specific detection of the CC and specificity for their target sequences, as well as enhanced CC beholves action of their target sequences, as well as enhanced CC biological activity. AB071213 to AB072214 and ABP48191 to AB97230 CC represent DNA target sequences and zinc finger partides which are given
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RESULT 169
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                                                     The present invention describes a zinc finger protein (I) that binds to CC a target site, comprising a first (FI), a second (F2), and a third (F3) CC zinc finger, ordered FI, F2, F3 from N-terminus to C-terminus, where the CC target site comprises, in 3'-5' direction, a first (S1), a second (S2), and a third (S3) target subsite. Also described are: (I) a polypeptide (II) comprising (I); (2) a polynucleotide (III) encoding (I) or (II); and (C3) designing (M) (I) involves selecting the F1 zinc finger such that it binds to the S1 target subsite, selecting the F2 zinc finger such that it binds to the S2 target subsite, and selecting the F3 zinc finger such that it binds to the S3 target subsite, thus designing (I) concerning the F3 zinc finger such that the binds to the S3 target subsite, thus designing (I) that binds to a target site. (I) is useful for recognition of triplet that binds to a target site. (I) is useful for recognition of triplet concerning the subsite shaving the nucleotide G in the 5'-most position of the Subsite. (I) is useful in the C1 therapeutics and plant engineering. (I), (II) or (III) is useful in the C2 therapeutic methods to modulate the expression of a target region within C3 subject, in diagnostic methods for sequence specific detection of the target nucleic acid in a sample, and in assays to determined the C3 phenotype and function of gene expression. (I) has improved affinity can be proceed affinity and septiment of the process and zinc finger peptides which are given in the process and zinc finger peptides which are given in the process and zinc finger peptides which are given in the process and zinc finger peptides which are given in the process and zinc finger peptides which are given in the process and zinc finger peptides which are given in the process and zinc finger peptides which are given in the process and zinc finger peptides which are given in the process and zinc finger peptides which are given in the process and zinc finger peptides which are given in the process and zinc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Zinc finger
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABQ72126 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SANG-) SANGAMO BIOSCIENCES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
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                                         exemplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 B CCAGGCCA 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein related oligonucleotide target SEQ ID NO:2424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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5.2e+02;
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Sequence 9 BP; 0 A; 3 C;

4 G;

2 T; 0 other;

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                                                                Query Match
Best Local :
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Best Local &
                                                     Matches
                                                                                                                                        The present invention describes a group of genes consisting of 100 genes which are highly expressed in human dendritic cells; a group of genes which are expressed at a higher frequency in human dendritic cells than in human monocytes; and a group of genes which are expressed at lower frequency in human dendritic cells than in human monocytes. Each group of genes are characterised in that CDNAs of these genes respectively have the base sequences of SEQ ID NO:10 to 200 (AAC74014 to AAC74113) and SEQ ID NO:201 to 300 (AAC7414 to AAC74213), each is continuous with the base sequence of "CATG-3" located most closely to the poly-A region. The sequence 5'-CATG-3' located most closely to the poly-A region. The sequence involvement of dendritic cells in the immune system and for the study and diagnosis of diseases in which dendritic cells play a significant role,
                                                                                                                                                                                                                                                                                                                                                                           Groups of genes expressed in human dendritic cells at a greater or lesser extent than in monocytes for investigation and diagnosis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; dendritic cell; monocyte; immune system; diagnosis; cancer; autoimmune disease; tumour; ss.
                                                                                                       Sequence 10
                                                                                                                               e.g. cancers and autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 9; 95pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                autoimmune disease and
                                                                                                                                                                                                                                                                                                                                                                                                                                                Hashimoto S,
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                                                 Similarity
8; Conserv
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                      GGCCAGGC 13
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                                                   Conservative
                                                                                                       BP; 1 A;
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100.0%; Pr
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                                                                                                       3 C; 4 G;
                                                                             40.0%;
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Pred. No.
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                                            DB 1; L;
No. 1.1e+02;
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RESULT 171 ABA06204

(UYJO ) UNIV JOHNS HOPKINS

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AAH64388
AAH64388
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AAHC
AXX
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Hume
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Home
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a human normal hepatocyte expression gene group comprising 200 genes in the human normal hepatocyte. The cDNA of each gene comprises one of 200 fully defined nucleotide sequences as given in the specification. The gene group and the cDNAs corresponding to each of the genes in the group are useful in the diagnosis and treatment of human hepatopathy. The present sequence is a cDNA corresponding to a gene expressed by normal human
                                     24-NOV-1999;
                                                              21-NOV-2000;
                                                                                         31-MAY-2001
                                                                                                                                          Homo sapiens
                                                                                                                                                                  Human; transcriptome; gene expression pattern; cancer; drug screening; cancer diagnosis; cell specific gene expression; ss.
                                                                                                                                                                                                         Human ubiquitously expressed transcriptome sequence SEQ ID NO: 1228.
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                                                                                                                                                                                                                                                             AAH64388
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 9; 26pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human normal hepatocyte expression gene group
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8; Conserv
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                                                               2000WO-US31922
                                                                                                                                                                                                                                   (first entry)
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                                     99US-0448480
                                                                                                                                                                                                                                                                                      cDNA; 10
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100.0%;
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RESULT 173
AAH64389
ID AAH643
XX AAH643
XX AAH643
XX Human
XX Human;
XW Human;
XW Cancer
XX Homo s
PN WO2001
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XX Homo s
YX Claim
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Best Local :
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                                                The present invention describes a method of identifying the type of cell in a sample, involving determining which of the sequences
AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce
                                                                                                                                                                                 New isolated polynucleotides, useful type, such as cancer cell, comprises particular cell types -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a method of identifying the type of cell in a sample, involving determining which of the sequences AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated polynucleotides, type, such as cancer cell, coparticular cell types -
cancer specific gene expression, standardise expression and restore the function of a diseased cell or tissue. The present sequence is one of the transcriptomes described in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; transcriptome; cancer diagnosis; cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer specific gene expression, standardise expression and restore t function of a diseased cell or tissue. The present sequence is one of the transcriptomes described in the exemplification of the invention.
                                                                                                                                                  Claim 13; Page 67; 94pp; English.
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                                                                                                                                                                                                                                                                                                                                                     24-NOV-1999;
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100.0%; Pr
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No. 1.1e+02;
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RESULT 174
ANH64390
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RESULT 175
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                                                                                                                                                                                                                                                                       Sequence 10 BP; 1 A; 3 C; 4 G; 2 T; 0 other;
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RESULT 176
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                     24-NOV-1999;
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                                                                                                                                 cancer
                                                                                                                                          Human;
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Best Local
           The present invention describes a method of identifying the type of cell in a sample, involving determining which of the sequences AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce cancer specific gene expression, standardise expression and restore the
                                                                                                                                                                                                                                                                                                                                                                                                     Human;
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AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce
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                                                                                                        Claim 13;
                                                                                                                                   type, such as cancer cell, particular cell types -
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                                                                                                        67; 94pp; English.
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RESULT 179
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Best Local Similarity
Matches 8; Conserva
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  ABV84947;
                   ABV84947 standard; cDNA; 10
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                                                                                                                                                                                                                                                                  New isolated polynucleotides, useful type, such as cancer cell, comprises
                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; transcriptome; gene expression pattern; cancer; drug screening; cancer diagnosis; cell specific gene expression; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 10 BP; 1 A;
                                                                                                                                       Sequence 10
                                                                                                                                                                                                                                       Claim 13; Page 67; 94pp; English.
                                                                                                                                                                                                                                                                                                                Velculescu VE,
                                                                                                                                                                                                                                                                                                                                                   24-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                        31-MAY-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human ubiquitously expressed transcriptome sequence SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH64394;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 the transcriptomes described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAH64394 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                  (UYJO ) UNIV JOHNS HOPKINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                               GGCCAGGC 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGCCAGGC 10
                                                                                                                                                                                                                                                           cell
                                                                                                                                      BP; 1 A; 3 C; 4 G; 2 T; 0 other;
                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                   99US-0448480
                                                                                                                                                                                                                                                                                                                Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             40.0%; sc.
100.0%; Pred. No.
100.0%; Mismatches
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                                                                                                           100.0%;
                                                                                                                     40.0%;
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                                                                                                           Score 8; 1; Pred. No.
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                                                                                                                                                                                                                                                                  for identifying specific cell transcriptomes expressed in
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o. 1.1e+02;
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o. 1.1e+02;
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ABV78373
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Matches
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The invention relates to SAGE (serial analysis of gene expression) tags Crepresenting groups of genes which are differentially expressed in human CC chronic hepatitis C (CH) liver tissue or hepatitis C-induced thepatocellular carcinoma (HCC) compared with normal human liver tissue. The SAGE tags of this invention consist of a sequence of 10 nucleotides CC clocated downstream of the 5'-CATG-3' sequence motif lying nearest to the CC located downstream of the 5'-CATG-3' sequence motif lying nearest to the CC located downstream of the 5'-CATG-3' sequence motif lying nearest to the CC located downstream of the 5'-CATG-3' sequence motif lying nearest to the CC located downstream of the sencoded by the genes can thus be used to analyse the CC pattern of gene expression in particular cell types. The invention also CC relates to proteins encoded by the genes expressed in chronic hepatitis C liver tissue or HCC, antibodies against these proteins, and inhibitors CC constitis C liver tissue or HCC droups of genes differentially expressed in chronic hepatitis C liver tissue or HCC may be used for the diagnosis and CC constitute, and antibodies against the gene products may be used in the development of drugs to treat chronic hepatitis C and/or HCC. Sequences ABV84891-ABV84990 are SAGE tags representing 100 genes which are highly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                        SAGE tag; serial analysis of gene expression; human; Th2 cell activated T cell; T lymphocyte; immune response; expression p
                                                                                                                                   Human Th2
                                                                                                                                                                                                    29-NOV-2002
                                                                                                                                                                                                                                                                    ABV78373;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-631294/68
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human multiple HCC highly expressed genes SAGE tag #757
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
disorder;
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8; Conserv
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                                                                                                                                cell SAGE tag, SEQ ID NO:84
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              in hepatocellular carcinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Page 31; 139pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 1 A; 3 C; 4 G; 2 T; 0 other;
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Conservative
                                                                                                                                                                                                    (first
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100.0%; Pred. No. 1.1e+02;
live 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to SAGE (serial analysis of gene expression) tags representing groups of genes which are expressed in activated human Thi and/or Th2 cells. The SAGE tags of this invention consist of a sequence of 10 nucleotides located downstream of the 5. CATG-3. sequence motif lying nearest to the polyA region of cDNAs derived from a variety of genes. These tags serve to uniquely identify each transcript and can thus be used to analyse the pattern of gene expression in particular cell types. The invention also relates to proteins encoded by the genes expressed in Th1 and/or Th2 cells, antibodies against these proteins, and inhibitors of the expression of groups of genes that are expressed in either or both the two cell types. Groups of genes expressed in Th1 and/or Th2 cell types and be used for the diagnosis and treatment of Th1 and Th2-related disorders. Sequences ABV78340-ABV78389 are SAGE tags crepresenting 50 genes which are most highly expressed in Th2 cells.
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Best Local S
Matches 8
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Human maturation/activation dendritic cell expression gene group
                                                             WPI; 2002-127070/17
                                                                                                                                                                                                                                      22-MAY-2000; 2000JP-0150562
                                                                                                                                                                                                                                                                                                    27-NOV-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             maturation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; maturation/activation dendritic cell expression gene; tag;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human maturation/activation dendritic cell expression gene tag
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                                                                                                                 (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             activation; dendritic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 1 A;
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100.0%; Pr
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cell;
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                                                                                                                                            The present invention describes a human maturation/activation dendritic cell (DC) expression gene group consisting of 100 genes which show the highest expression among the genes expressed in human maturation/activation DC also described are: (1) a protein expressed by the above human maturation/activation DC expression gene; (2) an antibody against the protein; and (3) an antagonist against the expression of each gene belonging to the above gene group. The gene group is useful for the treatment and the diagnosis of various human diseases related to human DC. ABL42627 to ABL42926 represent specifically claimed human invention.
                                                                                                            Sequence 10 BP; 1 A; 3 C; 4 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                 Claim 1; Page 9; 41pp; Japanese
   δ
                                    Similarity
8; Conserv
   GGCCAGGC 13
                                       Conservative
                                                        40.0%;
100.0%;
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). 1.1e+02;
                                                                        Length 10;
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RESULT 182
AAQ51821/c
                                                         Multiple drug resistance; mdr-1; ribozyme; membrane protein; liver; resistance; chemotherapeutic agent; colchicine; doxorubicin; colon; actinomycin D; vinblastine; small intestine; kidney; adrenal gland; adenocarcinoma; bowel; transformed phenotype; promyelocyric leukemia; human; chronic myelogenous leukemia; CML; follicular lymphoma;
human; chronic myelogenous leukemia; CML; follicular B-cell acute lymphocytic leukemia; breast cancer; co neuroblastoma; lung cancer; genetic drift; mutation;
                                                                                                                                                                                                                                                                                                            25-MAR-2003
26-MAY-1994
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                                                                                                                                                                                                                                                mdr-1 mRNA ribozyme cleavable nucleotide NT585.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard;
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GGCCAGGC 10

26-AUG-1992; 26-AUG-1992; 26-AUG-1992; 26-AUG-1992; 07-DEC-1992 19-JAN-1993 14-MAY-1992; 26-AUG-1992; 14-MAY-1992 13-MAY-1993; 25-NOV-1993 WO9323057-A1 Homo sapiens 92US-0936422. 92US-0936531. 92US-0936532. 92US-0987131. 93US-0006122. 92US-0882822. 92US-0882885. 92US-0936110. 9208-0936421 93WO-US04573

ଜି Thompson

(RIBO-)

RIBOZYME PHARM INC

93US-0008910

multiple New enzymatic RNA molecules (ribozymes) associated with tumours or mRNA express drug \ expressed 1 which cleave from gene enc

Claim

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RESULT 183
AAQ96792
ID AAQ967
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                  23-DEC-1994;
14-FEB-1994;
21-FEB-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The sequences given in AAQ51816-24 represent areas of the multiple drug resistance (mdr-1) mRNA which are accessible to the ribozyme of the invention. The mdr-1 gene encodes a 170 kD integral membrane protein which confers resistance to certain chemotherapeutic agents, such as colchicine, doxorubicin, actinomycin D and vinblastine. The
                                                      New non-pathogenic HIV-1 strain carrying a deletion in its nef or LTR region - can be used in a vaccine to inhibit/reduce productive infection in an individual by a pathogenic strain
                                                                                                                                                                                                                                   (AURE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                   17-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HIV-1; AIDS;
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                  Claim 13; Page 193; 301pp; English
                                                                                                                                                                                                                                                                                                                                                                                        14-FEB-1995;
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                                                                                                                                                  1995-293115/38
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunodeficiency
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                                                                                                                                                                                                                                   AUSTRALIAN RED CROSS SOC NSW DIV. MACFARLANE BURNET CENT MEDICAL.
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                94AU-0000284.
94AU-0003864.
94AU-0004002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    virus type
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o. 1.1e+02;
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Best Local S
Matches 8
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Best Local S
Matches 8
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14-FEB-1994;
21-FEB-1994;
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                                                                                                                                                                                                                                      Attenuation of pathogenic HIV-1 strain NL4-3 involves deletion of 1 or more decanucleotides (AAQ96406-Q97018) from the nef gene and/or 1 or more decanucleotides (AAQ97019-Q97166) from the LTR region; the sequence of AAQ96406 corresponds to nucleotides 1-10 of the nef gene (AAQ950141). The resulting avirulent HIV strains are still capable of inducing an immune response in humans, and enable the generation of therapeutic, diagnostic and targeting agents against HIV-1 infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New non-pathogenic HIV-1 strain carrying a deletion in its nef gene or LTR region - can be used in a vaccine to inhibit/reduce productive infection in an individual by a pathogenic strain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                             Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (AURE-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HIV-1; AIDS; attenuation; vaccine; nef
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HIV-1 NL4-3
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8; Conserva
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MACFARLANE
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                                                                                                                        Similarity
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                                              GGGCCAGG 12
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                                                                                                                                                                                             BP; 1 A; 2 C; 7 G; 0 U; 0 other;
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                                                                                              Conservative
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94AU-0003864.
94AU-0004002.
                                                                                                                                                                                                                                                                                                                                                                                                                                         193; 301pp; English.
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BURNET CENT MEDICAL.
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                                                                                                                                               40.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Deacon NJ,
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0; Mismatches
                                                                                                                     Score 8;
Pred. No.
                                                                                              Mismatches
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                                                                                                                     DB 1; Le:
b. 1.1e+02;
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5. 1.1e+02;
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                                                                                                                                            Length 10;
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RESULT 186
AAX54584/c
ID AAX54584 standard; DNA; 10 BP
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AAQ88580/c
                                                                                                                         Query Match
Best Local S
Matches 8
                                                                                                                                                                                                    A DNA chip was prepared for analysing sequences contained in a 1.3kb fragment of human mitochondrial DNA from the D-loop region, the most polymorphic region of human mitochondrial DNA. The chip comprised a set of 268 overlapping oligonucleotide probes (see AQ88421-Q88684) of varying length (9-14 nucleotides) with varying overlaps arranged in a 1cm x 1cm array. Each position in the sequence was represented by at least one probe (usually 2 or more). DNA was amplified from six human donors and then transcribed to give the 1.3kb RNA transcripts which were fragmented and hybridised to the chip. For each individual, a unique hybridisation fingerprint was produced on the chip; all differences could be correlated with differences in the cloned genomic DNA sequence.
                                                                                                                                                                            Sequence 10 BP; 1 A; 4 C; 5
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tiling strategy; immobilised nucleic acid probe array; mitochondrial DNA; D-loop region; biological chip; hybridisation fingerprint; interrogation position; ss.
                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 108;
                                                                                                                                                                                                                                                                                                                                                                                                          New arrays of oligo:nucleotide probes sequences with variants for detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-AUG-1994;
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8; Conserv
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Sheldon EL;
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                                                                                                                          Conservative
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93US-0143312.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "3'-end of probe is covalently attached
to chip surface"
                                                                                                                                     100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fodor SPA,
                                                                                                                                                                                                                                                                                                                                                                               223pp; English.
                                                                                                                   J. 0%;
0;
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Lobban PE, Miyada CG,
                                                                                                                         Score 8; DB 1; Pred. No. 1.1
0; Mismatches
                                                                                                                                                                            G; 0 U; 0 other;

    used for comparing known
of mutation(s) and sequencing.

                                                                                                                                    DB 1; Ler
5. 1.1e+02;
                                                                                                                                                Length 10;
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G, Morris MS;
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                                                                                                                                                               The specification describes antisense oligonucleotides (AAX52869-X55271) CC directed against at least 2 mRNAs selected from target genes, coding and conno-coding regions of RNAs corresponding to target genes, gene cc initiation codons, genomic flanking regions, intron-exon borders, the cc 5'-end, the 3'-end and the juxta-section between coding and non-coding cregions and all segments of RNAs encoding proteins associated with one cor more diseases, conditions or mixtures. The antisense oligonucleotides can be derived from sequences AAX55272-74. These multiple target coligonucleotides (specifically AAX55180-271) can be used for the can be derived from sequences and conditions. Typical diseases and conditions are those associated with impaired respiration and conditions are those associated with impaired respiration and conditions are those associated with impaired respiration, and the conditions are those associated with impaired respiration, and the conditions are those associated with impaired respiration, and largic rhinitis, acute asthma, allergies, asthma, impeded conditions are those associated with impaired respiration, construction, respiration, pulmonary vasoconstriction, emphysema, chronic constructive pulmonary disease (COPD), and cancers such as leukemias, corcinomas, carcinomas e.g. colon cancer, breast cancer, lung cancer, colon cancer, breast cancer, melanoma, concer, melanoma, melanoma, metastasses, as well as all types of cancers which may metastasize cor have metastasized to the lungs, including breast and prostate cancer.
                                                                               Best Loc
Matches
                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary hypertension; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer; hepatic cancer; hepatic cancer; colon carcer; breast cancer; lung cancer; melanoma; hepatic metastasis;
                                                                                                                                                             Sequence 10 BP; 0 A; 2 C; 6 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 45; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vasoconstriction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides used in treatment of, e.g. pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-229400/19
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17-SEP-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 prostate cancer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense oligonucleotide; multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vasoconstriction; inflammation; allergic rhinitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX54584;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO9913886-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human beta tryptase antisense oligonucleotide fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYEC-) UNIV
                                                                                                    Local
  ω
                                          ω
                                                                             Similarity
8; Conserv
                                          CCAGGCCA 15
CCAGGCCA 1
                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EAST CAROLINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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97US-0059160
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                                                                                                  100.0%;
                                                                                                                       40.0%;
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                                                                                                    Score 8;
Pred. No.
                                                                             Mismatches
                                                                                                    DB 1; Le
                                                                               0
                                                                                                                     Length 10;
                                                                               Indels
                                                                               0,
                                                                               Gaps
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RESULT 187

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RESULT 188
AAF20153/c
ID AAF201
XX
AC AAF201
XX
DT 14-MAR
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                                                                                                                                                                                                                                                                                                A method has been developed for the diagnosis of cancer in potentially complastic samples. The method comprises comparing the level of transcription between RNA transcripts in two tissue samples (which are of the same type), where the first sample is potentially neoplastic, and categorized as neoplastic if its level of transcription is lower than categorized as neoplastic if its level of transcription is lower than that of the second sample. The transcript is selected from Alu, RAS, CC under the constant of the second sample is selected from Alu, RAS, CC and LIMK-2, and it is identified by a tag selected from ribosomal protein L13a, alpha-tubulin (T1) and (T2), thymosin beta-4, and gamma-actin. The present sequence represents a serial analysis of gene expression (SACE) tag from the present invention. The use of SAGE tags provides an extensive profile of gene expression in rat embryo cc suppression gene. The discovery of new SAGE tags, which are regulated by p53, enables the diagnosis of genes that are related to cell cycle control and funcurearists.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX18625/c
                                                                                                                                                                                                Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Diagnosis of cancer in potentially neoplastic samples - by comparing the level of transcription between RNA transcripts in two tissue samples, useful for providing an extensive profile of gene expression in rat embryo fibroblast (REF) cells
 14-MAR-2001
                               AAF20153;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             p53; serial analysis of gene expression; SAGE tag; cancer; neoplastic; rat embryo fibroblast; REF; tumour suppressor; cell cycle control; tumourigenesis; diagnosis; ss.
                                                            AAF20153 standard; DNA; 10
                                                                                                                                                                                                                                                             Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Page 15; 32pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-106079/09
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-JUL-1997;
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                                                                                                                                                                                                                                                             BP; 1 A;
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 (first entry)
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                                                                                                                                                                                                                                                             3 C; 4 G;
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                                                                                                                                                                                                              Score 8; 1; Pred. No.
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                                                                                                                                                                                                                                                             2 T;
                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                             0 other;
                                                                                                                                                                                                                             DB 1;
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Human beta tryptase polynucleotide fragment #1720
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human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic; respiratory obstruction; pulmonary obstruction; impeded respiration; surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; cancer; ss. surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pain; cystic fibrosts; allergic rhinitis; pulmonary hypertension; emphysema; pulmonary transplantation rejection; chronic obstructive pulmonary disease; pulmonary infection; bronchitis; Low adenosine antisense oligonucleotide; phosphorothioate; allergy;

Homo sapiens

WO200062736-A2

24-MAR-2000; 2000WO-US08020

06-APR-1999; 9908-0127958

(UYEC-) UNIV EAST CAROLINA. (NYCE/) NYCE J W.

WPI; 2000-679539/66

Low adenosine (A) content antisense oligonucleotides which do not trigger adenosine receptors during metabolism, useful e.g. for treating cancers and respiratory obstructions -

Claim 14; Page 141; 1592pp; English.

immunoglobuling and antibodies, antibody receptors, cytokines and chemokines, endogenously produced specific and non-specific enzymes, binding proteins, adhesion molecules and their receptors, cytokine and chemokine receptors, adenosine receptors, bradykinin receptors, central nervous system (CNS) and peripheral nervous and non-nervous system receptors, CNS and peripheral nervous and non-nervous system peptide transmitters, defensins, growth factors, vasoactive peptides and receptors, binding proteins and malignancy associated proteins. The antisense oligonucleotides may be used in this way to treat disorders including respiratory obstruction (especially pulmonary obstruction and/or bunchoconstriction) and/or lung inflammation, allergy(ies) and/or surfactant hypoproduction which are associated with a disease or immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. The antisense oligonucleotides and (1) can be used to down-regulate the expression and or activity of target polypeptides associated with lung/respiratory disorders and malignancies, such as stimulating and activating peptide factors and transmitters, transcription factors, activating peptide factors and transmitters, transcription factors, The present invention describes low adenosine (A) content antisense oligonucleotides and compositions (I) comprising them. In the antisense oligonucleotides the A is replaced by a 'Universal' or alternative base. (I) can have respiratory, bronchodilator, antiinflammatory, analgesic, pulmonary transplantation rejection, pulmonary infections, bronchis and/or cancer. AAF18434 to AAF21543 represent human polynucleotide fragments and antisense oligonucleotides used in the exemplifications. condition selected from pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome (RDS), pain, cystic fibrosis (CF), allergic rhinitis (AR), pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD) the exemplification of

Sequence 10 BP; 0 A; 2 C; 6 G; 2 T; 0 other;

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Matches
          Best Local
                     Query Match
Similarity
8; Conserv
Conservative
         40.0%; Score 8; DB 1; Le
100.0%; Pred. No. 1.1e+02;
0,
Mismatches
                   Length 10;
Indels
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Gaps
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CCAGGCCA 15

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                                                                                                                                                                 most frequently in human monocytes. The cDNA of each gene has a gequence fully defined in the specification, and lacking the CATG sequence located adjacent to polyA region. Also described are:

(1) an antibody specifically for the protein encoded by any of the genes; (2) oligonucleotides obtained from the cDNA sequences;

(3) 380 human genes which are expressed most frequently in human macrophages, differentiated from human monocytes by granulocyte-macrophage colony-stimulating factor, the cDNA of each gene has a fully defined sequence, given in the specification, lacking the base sequence CATG located most closely to the poly A region;

(4) an antibody specifically for the protein encoded by any of the genes of (3); and (5) oligonucleotides obtained from the cDNA sequences of (3). The genes and cDNAs, are used for the study of gene specificity and disease onset mechanism e.g. oncogenesis, genetic diseases, drug clevelopment and diagnosis. AAA56107 to AAA56586 represent specifically connocytes and macrophages.
                                                            Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Genes most frequently expressed in human monocytes and GM-macrophages and M-macrophages studied and with cDNAs characterized, for study of gene specificity, disease onset mechanism, drug development and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            granulocyte-macrophage colony-stimulating factor; characterisation; GM-CSF; identification; diagnosis; gene specificity; oncogenesis; disease onset mechanism; genetic disease; drug development; ss.
                                                                                                                          Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes 100 human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hashimoto S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human macrophage gene Tag oligonucleotide sequence
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                                                              Local Similarity
les 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    43; Page 129; 138pp; Japanese.
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                            CAGGCCAA 16
                                                                                                                                                          and macrophages.
                                                                                                                          BP; 1
                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matsushima K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SCI & TECHNOLOGY CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98JP-0307532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99WO-JP05982
                                                                                                                          A.
                                                       100.0%;
                                                                                                                          2 C; 3
                                                                                              40.0%;
                                                          Score 8; DB 1; Pred. No. 1.1
0; Mismatches
                                                                                                                          G; 4 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Suzuki T;
                                                                                                                                                                                                                                                                                                                                                                                                                   human genes, which are expressed
The cDNA of each gene has a
                                                                                                                            other;
                                                                           DB 1; Le
). 1.1e+02;
                                                                                         Length 10;
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                                                            Gaps
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RESULT 190
AAA34031/c
The present invention describes a new composition comprising an antisense oligonucleotide (ON) with low adenosine (up to 15%), which targets nucleic acids involved in bronchoconstriction, allergies, and/or inflammation. The ON can have antiinflammatory, antiallergic, and can tiasthmatic, cytostatic and analgesic activities. The compositions are useful for the treatment of diseases associated with inflammation, fingularly lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating e.g. ischaemic conditions, pulmonary vasoconstriction, allergies, fibrosis, pulmonary hypertension, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukaemias, lymphomas, carcinomas, and cancers which may metastasise to the lungs, including breast and prostate cancer. The reduction of the adenosine content of the ONs reduces side effects. The A-containing ONs break down with the release of deoxyadenosine which activates adenosine receptors causing bronchoconstriction and inflammation. AAA33311 to AAA33312 represent the
                                                         nucleotide sequences given in the sequence listing from the present invention, which correspond to SEQ ID NO:1 to 2815, and then the last 185 sequences are also called SEQ ID NO:1 to 185, but the sequences differ from the previously named sequences. SEQ ID NO:11 to 1680 (AAA32323 to AAA33992) are specifically claimed ONs from the present invention. N.B. Sequences given in the disclosure of the present invention do not match up with their corresponding SEQ ID NO: sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstriction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 478; 1343pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-205971/18.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer; leukaemia; lymphoma; carcinoma; metastasis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           pulmonary hypertension; chronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; lung disease; ischaemic condition; pulmonary vasoconstriction; asthm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  phosphorothioate; impaired respiration; inflammation; allergy;
allergic disease; bronchoconstriction; inhibitor; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human adenosine receptor related polynucleotide SEQ ID NO:1720.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               respiratory distress syndrome; pain; cystic fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                         listing.
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obstructive pulmonary disease; COPD;
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Query Match
Best Local Similarity
Matches 8; Conserv

Conservative

40.0%; but 100.0%; Pr

Score 8; DB 1; Pred. No. 1.1
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DB 1; Le o. 1.1e+02;

Length 10;

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Gaps

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Sequence 10

BP; 0 A; 2 C; 6 G;

2 T; 0 other;

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CCAGGCCA 15

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RESULT 191
AAZ77634/c
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19-JUN-1998;
19-JUN-1998;
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19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SAGE tag; serial analysis of gene expression; antigen-presenting cell; APC; monocyte-derived dendritic cell; differential gene expression; immunostimulatory cofactor; costimulatory factor; CTL;
                Claim 1; Page 65; 130pp; English
                                                                                                 (GENZ )
(ROBE/)
(SHAN/)
                                                                                                                                                                                             19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cytotoxic T-lymphocyte; tumour antigen; immunotherapy; anticancer; ss
                                                              WPI; 2000-106077/09.
                                                                                                                                                19-JUN-1998;
19-JUN-1998;
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19-JUN-1998;
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                                                                                                                                                                  19-JUN-1998;
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                                                                                                 ) GENZYME CORP.
) ROBERTS B L.
) SHANKARA S.
                                  ed polynucleotides differentially expressed in antigen-presenting useful in gene vaccines against cancer -
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                                                                                ВЪ,
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                                                                                Shankara
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98US-0090036
98US-0090039
98US-0090040
98US-0090041
98US-0090042
98US-0090042
98US-0090043
98US-0090045
98US-0090045
98US-0090075
98US-0090072
98US-0090077
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98US-0090078
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98US-0089994.
98US-0089997.
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98US-0089992.
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98US-0089878.
                                                                                                                                                                                                                                                                                                                                                                                                     98US-0089833
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                                  gene vaccines against
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Sequences AAZ77573-Z79709 represent SAGE (serial analysis of gene

19-JUN-1998; 19-JUN-1998; 19-JUN-1998; 19-JUN-1998; 19-JUN-1998; 19-JUN-1998; 19-JUN-1998;

98US-0089833. 98US-0089844. 98US-0089878. 98US-0089991. 98US-0089991. 98US-0089993.

18-JUN-1999;

99WO-US13800

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RESULT 192
AAZ79103/C
ID AAZ791
XX AAZ791
XX 10-APR
XX SAGE Human
XX SAGE C
KW APC; m
KW immuno
XW cytoto
XX Homo s
XX W09965
XX W09965
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PR 19-JUN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC activation of the cycotoxic immune response, particularly against tumour cells. Tumour antigen presentation via the MHC (major histocompatibility C complex) and subsequent recognition by T-cell receptors is alone CC insufficient to activate a robust cyrotoxic immune response that can CC lyse the tumour cells, immunostimulatory cofactors also being required CC for efficient activation of cytotoxic T-lymphocytes (CTLs). Nucleic acid CC sequences identified using the SAGE tags have several potential uses. CT They may be used in vaccines to induce an immune response, particularly against a tumour antigen; to modulate the genotype of an APC; to screen CC for agents that modulate expression of differentially expressed genes in CC diagnosis, prognosis and monitoring of diseases related to abnormal CC differentially expressed genes, or of their encoded proteins, can be used to identify cells as belonging to the monocyte lineage. Cells containing these genes, or of their encoded proteins, can be used to identify cells as belonging to the monocyte lineage. Cells containing these genes and apc-associated costimulatory factors ensures adequate presentation of a population of antigen-specific effector cells) and vectors containing them are used in gene therapy. (Or -daministration of the antigen presentation of co-estimulatory signals, migration to T cell-rich sites, cecretion of T cell growth factors and secretion of chemokines for the recruitment of immune effector cells.
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SAGE tag; serial analysis of gene expression; antigen-presenting ce APC; monocyte-derived dendritic cell; differential gene expression; immunostimulatory cofactor; costimulatory factor; CTL; cytotoxic T-lymphocyte; tumour antigen; immunotherapy; anticancer;
                                                                                                                                                                                                                                                                                                                                        23-DEC-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            expression) tags used to identify mRNA transcripts encoding immunostimulatory cofactor proteins which are preferentially or differentially expressed in monocyte-derived dendritic cells compared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human dendritic cell SAGE tag,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (APC) -associated costimulatory factors play an important role in the
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100.0%;
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0; Mismatches
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lo. 1.1e+02;
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19-JUN-1998;
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                             differentially expressed genes, or of their encoded proteins, can be used to identify cells as belonging to the monocyte lineage. Cells containing these genes can be used in active immunotherapy (or to stimulate production of a population of antigen-specific effector cells) and vectors containing them are used in gene therapy. Co-administration of tumour antigens and APC-associated costimulatory factors ensures adequate antigen presentation to endogenous APCs and upregulates the APCs for the presentation of co-stimulatory signals, migration to T cell-rich sites, secretion of T cell growth factors and secretion of chemokines for recruitment of immune effector cells.
                                                                                                                                                                                                                    complex) and subsequent recognition by T-cell receptors is alone insufficient to activate a robust cytotoxic immune response that can lyse the tumour cells, immunostimulatory cofactors also being required for efficient activation of cytotoxic T-lymphocytes (CTLs). Nucleic acid sequences identified using the SAGE tags have several potential uses. They may be used in vaccines to induce an immune response, particularly against a tumour antigen; to modulate the genotype of an APC; to screen for agents that modulate expression of differentially expressed genes in an APC; and as hybridisation probes/amplification primers for the diagnosis, prognosis and monitoring of diseases related to abnormal expression of these genes. Detection of their accordance in the differentially expressed genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                      (APC)-associated costimulatory factors play an important role in the activation of the cytotoxic immune response, particularly against tumour cells. Tumour antigen presentation via the MHC (major histocompatibility
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       differentially expressed in monocyte-derived dendritic cells compared with monocytes. Some of the transcripts correspond to known genes or ESTs (expressed sequence tags) which were previously unknown to be preferentially or differentially expressed in dendritic cells, while other transcripts correspond to novel genes. Antigen-presenting cell other transcripts correspond to novel genes. Antigen-presenting cell
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         expression) tags used to identify mRNA transcripts encoding immunostimulatory cofactor proteins which are preferentially
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      useful in gene vaccines against cancer
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) ROBERTS B L.
) SHANKARA S.
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98US-0089997.
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Query Match Best Local Similarity

100.0%; 1 C; 4

Score 8; DB 1; Le Pred. No. 1.1e+02;

Length 10;

Sequence 10

N Α;

G; 3 T; 0 other;

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RESULT 193
AAZ79322/c
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                      Claim 1; Page 115; 130pp; English
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                                                                cells, useful in gene vaccines against
                                                                  Isolated polynucleotides differentially expressed in antigen-presenting cells, useful in gene vaccines against cancer -
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(ROBE/)
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19-JUN-1998;
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ROBERTS E
SHANKARA
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B L.
A S.
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RESULT 194
AAZ81594/c
ID AAZ815
XX AAZ815
XX AAZ815
XX O7-APR
XX Metast
XX Human;
KW non-me
KW non-me
KW non-me
XX Homo s
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19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  differentially expressed genes, or of their encoded proteins, can be used to identify cells as belonging to the monocyte lineage. Cells containing these genes can be used in active immunotherapy (or stimulate production of a population of antigen-specific effector cells) and vectors containing them are used in gene therapy. Co-administration of cumour antigens and APC-associated costimulatory factors ensures adequate antigen presentation to endogenous APCs and upregulates the APCs for the presentation of co-stimulatory signals, migration to T cell-rich sites, secretion of T cell growth factors and secretion of chemokines for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    They may be used in vaccines to induce an immune response, particularl against a tumour antigen; to modulate the genotype of an APC; to scree for agents that modulate expression of differentially expressed genes an APC; and as hybridisation probes/amplification primers for the diagnosis, prognosis and monitoring of diseases related to abnormal expression of these genes. Detection of the dendritic cell
                                                                                                                                                                                                                                                                                                                                                                                                                    23-DEC-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antimetastatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; metastatic breast tumour tissue; breast cancer; tag; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Metastatic breast tumour cell upregulated transcript tag #828.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-APR-2000
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                                                                                                                                                                                                                                                                                                                                    18-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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        (GENZ ) GENZYME CORP
                                                                                        19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequences AAZ77573-Z79709 represent SAGE (serial analysis of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
8; Conserv
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                                                                                98US-0089853.
98US-0089997.
98US-0090039.
98US-0090040.
98US-0090041.
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                                                                                                                                                                                                                                                                                                                                    99WO-US13647
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vaccine; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4 C; 4 G; 1 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   40.0%;
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ne; diagnosis; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells).

CC AAX83942 to AAX86677 represent tags corresponding to distinct transcripts (c) that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic. CC Diagnosis is by standard immunoassays or hybridisation/amplification (c) treatment of breast cancer particularly where metastatic monitoring and treatment of breast cancer, particularly where metastatic. CC Diagnosis is by standard immunoassays or hybridisation/amplification (c) promoters from the transcripts are used to direct expression, in selected continuous of e.g. therapeutic genes (also ribozymes or antisense (c) sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; (bor diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be used to be anticent and for inclusions.
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19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
                                                                                                                          23-DEC-1999
                                                                                                                                                        WO9965928-A2
                                                                                                                                                                                                                                 Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer;
                                                                                                                                                                                                                                                                                 Metastatic
                                                                                                                                                                                                                                                                                                              07-APR-2000
                                                                                                                                                                                                                                                                                                                                                                           AAZ82082 standard; DNA; 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis, prevention and treatment of cancer -
                                                                                           18-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                adoptive
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CGGGCCAG 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cells, e.g. cy immunotherapy.
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                                                                                                                                                                                                                                                                               breast tumour cell upregulated transcript tag #1316
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           isolate populations of educated,
              98US-0089853.
98US-0089997.
98US-0090039.
98US-0090040.
                                                                                           99WO-US13647
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100.0%;
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Pred. No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T lymphocytes,
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Best Local S
Matches 8
 19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in select cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences). particularly an antigen-encoding sequence for use in gene ocell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines, for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be use to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cyrotoxic T lymphocytes, and these used for
                                                                                                                                                                                        Human; metastatic breast tumour tissue; breast cancer; tag; non-metastatic breast tumour tissue; gene therapy; anticance antimetastatic; vaccine; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     breast tumour tissue (i.e. are downregulated in metastatic breast cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where meta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells).
AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast times.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis, prevention and treatment of cancer -
                                                                                                                                                                                                                                                          Metastatic
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                                                                                                                               WO9965928-A2
                                                                                                                                                             Homo sapiens
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                                                                18-JUN-1999;
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(ROBE/)
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) ROBERTS B L.
) SHANKARA S.
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8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                   GGGCCAGG 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunotherapy
                                                                                                                                                                                                                                                        breast tumour cell upregulated transcript
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98US-0089853.
98US-0089997.
98US-0090039.
                                                                99WO-US13647.
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Pred. No.
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5. 1.1e
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                       tag #2298
                                                                                                                                                                                                                         primer;
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reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in select sequences, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene cell-based vaccines, for diagnosing breast cancer and for raising useful in vaccines, for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be use to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for
                                                                                                                                                                                                                                                                               AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GENZ )
(ROBE/)
(SHAN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     prevention and treatment of cancer
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19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolated polynucleotides differentially and non-metastatic breast cancer cells,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ) GENZYME CORP.
) ROBERTS B L.
) SHANKARA S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BL,
                                      immunotherapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 121; 219pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shankara
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98US-0090041.
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Sequence 10 BP; 1 A; 2 C; 5 G; 2 T; 0 other;

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                       Query Match
Best Local S
Matches 8
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           v
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GGGCCAGG 10
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                        Mismatches
                     No. 1.1e+02;
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RESULT 197
AAZ83312/C
ID AAZ833
XX AAZ833
XX O7-APR
XX HUMAN,
KW HOMAN,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Metastatic breast tumour cell upregulated transcript tag #2546.
19-JUN-1998;
                                                                                                             18-JUN-1999;
                                                                                                                                                                                                                          23-DEC-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antimetastatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; metastatic breast tumour tissue; breast cancer; tag; primer;
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98US-0089853
                                                                                                             99WO-US13647
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tumour tissue; gene
ne; diagnosis; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          therapy; anticancer;
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RESULT 198
AAZ83787/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       promoters from the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis, prevention and treatment of cancer -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   monitoring and treatment of breast cancer, particularly where metastable possession is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while
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(ROBE/)
                                                                                                                          antimetastatic;
                                                                                                                                                       Human; metastatic breast tumour tissue; breast cancer; tag; primer;
                                                                                                                                                                                        Metastatic breast
                                                                                                                                                                                                                      07-APR-2000
                                                                                                                                                                                                                                                     AAZ83787;
                                                                                                                                                                                                                                                                                   AAZ83787 standard; DNA; 10
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18-JUN-1999;
                             23-DEC-1999.
                                                                                                                                          non-metastatic
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98US-0090040.
98US-0090041.
                                                                                                                          breast tumour tissue; gene therapy; vaccine; diagnosis; ss.
99WO-US13647.
                                                                                                                                                                                   tumour cell upregulated transcript tag #3021
                                                                                                                                                                                                                                                                                                                                                                                                                       40.0%; 5r
100.0%; Prr
                                                                                                                                                                                                                    entry)
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0; Mismatches
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o. 1.1e+02;
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Matches 8; Conserv
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19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
                                                                                             Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer;
                                                                                                                                                                   07-APR-2000
                                                                                                                                                                                                                         AAZ83855 standard; DNA; 10
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Isolated polynucleotides differentially and non-metastatic breast cancer cells, prevention and treatment of cancer -
                                                    Homo sapiens
                                                                                antimetastatic;
                                                                                                                                       Metastatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     transcripts that
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(ROBE/)
(SHAN/)
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ROBERTS
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                                                                                                                                                                                                                                                                                             GGGCCAGG
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                                                                                                                                       breast
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98US-0089997.
98US-0090039.
98US-0090040.
98US-0090041.
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                                                                                vaccine;
                                                                                                                                       tumour
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                                                                                diagnosis;
                                                                                                                                       cell
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                                                                                                                                      upregulated transcript tag
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                                                                                 ; gene
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                                                                                                                                                                                                                                                                                                                                                                                                          other;
                                                                                                                                                                                                                                                                                                                                                                DB 1;
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23-DEC-1999

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RESULT 200
AAZ83864
ID AAZ838
XX AAZ838
XX AZ838
XX O7-APR
XX OF Metast
XX Metast
XX Human;
KW Human;
KW Antime
XW Antime
XX ANTime
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic
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                                                                                                                      Metastatic breast tumour cell upregulated transcript tag #3098
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                                 Homo sapiens
                                                               antimetastatic;
                                                                            non-metastatic breast
                                                                                          Human; metastatic breast tumour tissue; breast cancer; tag; primer;
                                                                                                                                                                                        AAZ83864;
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(ROBE/)
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19-JUN-1998;
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19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                             Conservative
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98US-0090040.
98US-0090041.
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                                                              breast tumour tissue; gene therapy;
vaccine; diagnosis; ss.
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100.0%;
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Pred. No.
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RESULT 201
AAZ84284
ID AAZ842
XX
AC AAZ842
XX
DT 07-APF
DT 07-APF
XX
Metast
XX
KW Human;
KW non-mc
KW antime
XX
OS Homo:
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Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer; antimetastatic; vaccine; diagnosis; ss.

Homo sapiens

Metastatic 07-APR-2000 AAZ84284

breast tumour

cell downregulated transcript

#3518.

(first entry)

AAZ84284 standard; DNA; 10

ВP

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CAGGCCAA 16

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Matches
                                                            Best
                                                                                                   Query Match
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19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
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AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 promoters from the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides can be used therapeutic agents. Host cells that produce the polypeptides can be used
                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                 adoptive
                                                                                                                                                                                                                                                                                                                                                                        to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for % \left( 1\right) =\left\{ 1\right\} =\left\{
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(ROBE/)
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ROBERTS E
SHANKARA
     Similarity
8; Conserv
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     Conservative
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98US-0090041
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98US-0090039
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                                                                                                                                                                                                                        G; 0 U; 0 other;
                                                       Score 8; 1
Pred. No.
                Mismatches
                                                       DB 1; Ler
No. 1.1e+02;
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Gaps
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AAZ85316/C
ID AAZ853
XX AZ853
AC AAZ853
XX O7-APR
XX XX V7-APR
XX MCLASE
XX MCLASE
XX HUMMAR;
XW NON-Me
KW ANTIME
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Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer; antimetastatic; vaccine; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis,
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19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
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                                                                  Metastatic
                                                                                                                                 AAZ85316
                                                                                                                                                                 AAZ85316 standard; DNA; 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              prevention and treatment of
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ROBERTS B L.

SHANKARA S.
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                                                                                                                                                                                                                                               GGGCCAGG
                                                                breast
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                                                                                               (first entry)
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98US-0090039
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98US-0090041
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                                                             tumour cell downregulated transcript tag #4550.
                                                                                                                                                                                                                                                                                                         100.0%; F1
                                                                                                                                                                                                                                                                                                           40.0%; Score 8; DB 1
100.0%; Pred. No. 1.1
tive 0; Mismatches
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No. 1.1e+02;
0;
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RESULT 203
AAZ85453/c
ID AAZ854
XX AAZ854
AC AAZ854
XX O7-APR
XX Metast
XX Metast
XX Human;
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                           that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as specific antibodies (Ab). Ab are used to detect the polypeptides or as the reactions are used to detect the polypeptides or as the reactions.
                                Metastatic
                                                                07-APR-2000
                                                                                               AAZ85453
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                                                                                                                                                                                                                                                                                                                                                                   adoptive
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19-JUN-1998;
19-JUN-1998;
                                                                                                                              AAZ85453
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GENZ )
(ROBE/)
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ROBERTS B L.
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8; Conserv
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                                                                                                                              standard; DNA;
                                                                                                                                                                                                                                        GCCAGGCC 14
                                                                                                                                                                                                                                                                                                                                                                   immunotherapy.
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                                                                                                                                                                                                            GCCAGGCC
                                breast tumour cell downregulated transcript tag
                                                                                                                                                                                                                                                                                                                                      BP; 1 A; 4 C;
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                                                              (first entry)
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98US-0090040
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Jo. 1.1e+02;
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Human; metastatic breast tumour tissue; breast cancer; tag; primer;

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RESULT 204
AAZ85585/c
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Best Local Similarity
---hes 8; Conserv:
      DX SX FX B
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19-JUN-1998;
19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis, prevention and treatment of cancer -
Metastatic breast tumour cell downregulated transcript tag #4819
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                monitoring and treatment of breast cancer, particularly where metastatic Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while
                                 07-APR-2000
                                                                AAZ85585
                                                                                              AAZ85585 standard; DNA; 10 BP
                                                                                                                                                                                                                                                                                                Sequence 10
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(ROBE/)
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antimetastatic; vaccine; diagnosis; ss.
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                                                                                                                                                                                                                                                                                                                                           cells,
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                                                                                                                                                                                                                                     Conservative
                                 (first entry)
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98US-0089997.
98US-0090039.
98US-0090040.
98US-0090041.
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0; Mismatches
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RESULT 205
AAZ85596/c
ID AAZ855
XX
AC AAZ855
XX
AC AAZ855
XX
DT 07-APR

AA285596 standard; DNA; 10

ВР

07-APR-2000 AAZ85596;

(first entry)

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GGGCCAGG 12

Query Match Best Local S Matches 8

l Similarity 8; Conserva

40.0%;

100.0%;

Score 8; DB 1; Pred. No. 1.1
0; Mismatches

DB 1; Le o. 1.1e+02;

Length 10; Indels

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Conservative

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AAZ80767 to AAZ83941 represent tags corresponding to distinct CC transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, compitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification creations. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected concerns, of e.g. therapeutic genes (also ribozymes or antisense concernses), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines, for diagnosing breast cancer and for raising
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19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
Sequence 10
                                                                                 specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be use to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 187; 219pp; English.
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(ROBE/)
(SHAN/)
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                                                           adoptive
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) ROBERTS B L
) SHANKARA S.
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                                                           immunotherapy.
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   other;
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ne therapy; anticancer;
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  AAZ85958;
                                        AAZ85958 standard; DNA; 10
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(ROBE/)
(SHAN/)
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ROBERTS B L.
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98US-0090041.
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RESULT 207 ABA06090 ID ABA060

ABA06090 standard; cDNA; 10 BP

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                                                                                                                                                                adoptive immunotherapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Page 196; 219pp; English.
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                                                                                 Similarity
                     GGGCCAGG 12
                                                               40.0%; Scilarity 100.0%; F
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                                                               Score 8; DB 1; Pred. No. 1.1
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o. 1.1e+02;
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RESULT 208
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                                                                                                                                                                          Human ubiquitously expressed transcriptome sequence SEQ
 Velculescu
                                                             21-NOV-2000;
                                                                                 31-MAY-2001.
                                                                                                                         Homo sapiens
                                                                                                                                                        Human;
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                                        24-NOV-1999;
                                                                                                      WO200138577-A2
                                                                                                                                                                                                                                         AAH63781 standard; cDNA;
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                    (UYJO ) UNIV JOHNS HOPKINS
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diagnosis; cel
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8; Conserv
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VΕ,
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                                                             2000WO-US31922
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                                        99US-0448480
Vogelstein B,
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                                                                  The present invention describes a method of identifying the type of cell in a sample, involving determining which of the sequences
AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a method of identifying the type of cell in a sample, involving determining which of the sequences AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce
                cancer specific gene expression, standardise expression and restore the function of a diseased cell or tissue. The present sequence is one of the transcriptomes described in the exemplification of the invention.
                                                                                                                                                                                                                          New isolated polynucleotides, useful type, such as cancer cell, comprises
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; transcriptome; gene expression pattern; cancer diagnosis; cell specific gene expression;
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particular cell
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                                                                                                                                                                          59; 94pp; English
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RESULT 211
AAH64374/c
ID AAH643
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AC AAH643
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DT 20-SEP
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AAH64037/c
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Best Local Similarity
Matches 8; Conserv
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Best Local
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                                                                                                                                                                                                                                                      New isolated polynucleotides, useful type, such as cancer cell, comprises particular cell types -
 20-SEP-2001
                 AAH64374;
                                 AAH64374 standard; cDNA; 10
                                                                                                                                           Sequence 10 BP; 1 A; 4 C; 4 G; 1 T; 0 other;
                                                                                                                                                                                                                                     Claim 13; Page 59; 94pp; English.
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                                                                          GCCAGGCC 3
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(first entry)
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100.0%; Pr
                                                                                                        100.0%; F1
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                                                                                                          Score 8; DB 1; Pred. No. 1.1
0; Mismatches
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                                                                                                       DB 1; NO. 1.1e+02; 0;
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b. 1.le+02;
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RESULT 212
AAH64375/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated polynucleotides, useful type, such as cancer cell, comprises particular cell types -
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cancer
24-NOV-1999;
                                          21-NOV-2000; 2000WO-US31922
                                                                                    31-MAY-2001
                                                                                                                            WO200138577-A2
                                                                                                                                                                                                            Human; transcriptome; gene expression pattern; cancer; cancer diagnosis; cell specific gene expression; ss.
                                                                                                                                                                                                                                                                             Human ubiquitously expressed transcriptome sequence SEQ ID NO: 1215.
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                                                                                                                                                                                                                                                                                                                                                                   AAH64375;
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8; Conserv
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99US-0448480
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0; Mismatches
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o. 1.1e+02;
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RESULT 213
AAH32825
ID AAH328
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XX JP2001
XX GRA-APR
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Best Local S
Matches 8
                      The present invention describes an lipopolysaccharide (LPS) activated human monocyte expression gene group consisting of the high-ranking 50 genes of the high-tenking 50 genes of (AH32628 to AH32677) continuous to the base sequence of (AH32628 to AH32677) continuous to the base sequence 5. CATG-3. nearest to the polyA region. The gene group is useful for the development of new means for the diagnosis and the treatment of various human diseases in which human monocyte plays an important role.

AH32628 to AH32943 represent specifically claimed LPS activated human monocyte expression gene cDNA tags from the present invention. AH33944 represents an LPS activated human monocyte expression gene cDNA sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           LPS activated human monocyte expression gene
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RESULT 214
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                                                                                                                                                                                                                                                                                                               The present invention describes an lipopolysaccharide (LPS) activated human monocyte expression gene group consisting of the high-ranking 50 genes of the highest expression among the genes expressed by human monocyte stimulated by LPS in which the cDNA of each gene has the base sequence of (AAH32628 to AAH32677) continuous to the base sequence 5'-CATG-3' nearest to the polyA region. The gene group is useful for the development of new means for the diagnosis and the treatment of various human diseases in which human monocyte plays an important role. AAH32628 to AAH32943 represent specifically claimed LPS activated human monocyte expression gene cDNA tags from the present invention. AAH3294 represents an LPS activated human monocyte expression gene cDNA sequence encoding AAB98009, which are given in the exemplification of the present
                                                                                                                                                                                                                                Sequence 10
                                                                                                                                                                                                                                                                                          encoding AAB98009, invention.
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                                                                                                                   Conservative
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RESULT 215 AAF39487/c ID AAF394 XX

AAF39487 standard;

DNA;

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AAF39487;

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                                                                                                                                                          CC coding sequence of a yeast gene selected from a group of 745 NORF (not coding sequence of a yeast gene selected from a group of 745 NORF (not comprising a SAGE (serial analysis of gene expression) tag. Also CC comprising a SAGE (serial analysis of gene expression) tag. Also CC described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at CC least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate CC expression varies as in M1, where a test substance with a CC expression varies as in M1, where a test substance which modifies the CC expression of the yeast gene is a candidate antifungal drug; (3) a method CC (M3) for identifying human genes which are involved in cell cycle progression comprising contacting human DNA with a probe which comprises CC as in M1; and (4) a method (M4) for identifying a candidate drug as a candidate and (4) a method (M4) for identifying a candidate drug as a candidate drug and monitoring expression in the yeast cell with a CC candidate drug and monitoring expression in the yeast cell with a CC candidate drug and monitoring expression in the yeast cell of at least 1 (Contact of the cell cycle. The methods may be used to identify candidate drugs which cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The methods may be used as markers of phases of the cell cycle and for identification of antifungal drug. CC APF31368 to APF4064 represent SAGE tags used in the exemplification of the present invention. APF31262 to APF31367 represent linkers and PCR conversion in the SAGE method, in the exemplification of the present code in th
                                                                            Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yeast gene coding sequences comprising of gene expression (SAGE) tags, useful affecting phases of the cell cycle -
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                                                                                                                                                        Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example; Page 222; 419pp; English.
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o. 1.1e+02;
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ID AAF414
AC Yeast
CO Saccha
AC Saccha
AC WW Serial
AC WW Serial
AC AC UVJO
PF 14-JUN
AC COMPT
CC CO
                                                                                                   Cycle comprising administering a NORF gene whose expression varies by at CC cycle comprising and (2)/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a CC quast cell; and (b) monitoring expression of a NORF gene whose comprising contacting a test substance with a CC yeast cell; and (b) monitoring expression of a NORF gene whose comprising contacting a test substance which a CC quast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle (M3) for identifying human genes which are involved in cell cycle (M3) for identifying contacting human DNA with a probe which comprises CC at least 10 contiguous nucleotides of a NORF gene whose expression varies at 10 mM; and (4) a method (M4) for identifying a candidate drug as a CC expression in a yeast cell comprising contacting a yeast cell with a CC candidate drug and monitoring expression in the yeast cell with a CC candidate drug and monitoring expression in the yeast cell of at least 1 CNRF gene whose expression is affected by the class of drugs. The NORF CC genes may be used to identify candidate drugs which cell cycle. The methods may be used as markers of phases of the cell cycle. The methods may be used an intermination of phases of the cell cycle and for identification of antifungal drugs.

CC AAF33268 to AAF44064 represent SAGE method, in the exemplification of the present invention. AAF33267 to AAF33267 represent linkers and PCR invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             previously assigned open resource from a SAGE (serial analysis of gene expression) tag. Also comprising a SAGE (serial analysis of using NORF genes to affect the cell described are: (1) a method (M1) of using NORF gene whose expression varies by a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Yeast gene coding sequences comprising NORF genes with serial analy of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ζ,
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otated ORF; SAGE;
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RESULT 218
ABV84789
ID ABV847
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ABV84307
ID ABV843
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                                                                                                                                                                                                                                                                                                                         The invention relates to SAGE (serial analysis of gene expression) tags CC representing groups of genes which are differentially expressed in human CC chronic hepatitis C (CH) liver tissue or hepatitis C -induced CC hepatocellular carcinoma (HCC) compared with normal human liver tissue. The SAGE tags of this invention consist of a sequence of 10 nucleotides CC located downstream of the 5'-CATG-3' sequence motif lying nearest to the polyA region of cDNAs derived from a variety of genes. These tags serve to uniquely identify each transcript and can thus be used to analyse the pattern of gene expression in particular cell types. The invention also relates to proteins encoded by the genes expressed in chronic hepatitis C liver tissue or HCC, antibodies against these proteins, and inhibitors of the expression of groups of genes that are overexpressed in chronic hepatitis C liver tissue or HCC. Groups of genes differentially expressed in chronic hepatitis C tiver tissue or HCC may be used for the diagnosis and treatment of these diseases. Such genes, inhibitors of their expression of cor activity, and antibodies against the gene products may be used in the development of drugs to treat chronic hepatitis C and/or HCC. Sequences
                                                                                                                                                                Best Local S
Matches 8
ABV84789 standard; cDNA; 10
                                                                                                                                                                                                                                                                        ABV84291-ABV84390 are SAGE tags representing the 100 least highly expressed genes out of those genes which are underexpressed in ch hepatitis C liver tissue compared with normal liver tissue.
                                                                                                                                                                                                                                        Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 10; Page 13; 139pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human chronic hepatitis C tissue expression exasperating gene group comprises 100 high-ranking genes -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            expression pattern; differential expression; EST; expressed sequence tag;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SAGE tag; serial analysis of gene expression; human; chronic hepatitis CH; liver tissue; hepatocellular carcinoma; cancer; tumour; HCC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human chronic hepatitis C tissue underexpressed EST SAGE tag #117.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABV84307;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-JAN-2001; 2001JP-0012328
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                                                                                                                            10 AGGCCAAA 17
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                                                                                                                                                                Similarity 100
8; Conservative
                                                                                                                                                                                                                                    BP; 5
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 BP.
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                                                                                                                                                                                 Score 8;
Pred. No.
                                                                                                                                                                                                                                        other;
                                                                                                                                                                                 DB 1; Length 10; . 1.1e+02;
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RESULT 219
ABV78459
ID ABV784
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AC ABV784
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DT 29-NOV
XX
KW SAGE t
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                                                                                                                                                                                                                                                                                                                                                                                          CC representing groups of genes which are differentially expressed in human CC chronic hepatitis C (CH) liver tissue or hepatitis C-induced CC hepatocellular carcinoma (MCC) compared with normal human liver tissue. CC The SAGE tags of this invention consist of a sequence of 10 nucleotides CC located downstream of the 5'-CATG-3' sequence motif lying nearest to the CC polyA region of cDNAs derived from a variety of genes. These tags serve to uniquely identify each transcript and can thus be used to analyse the CC pattern of gene expression in particular cell types. The invention also crelates to proteins encoded by the genes expressed in chronic hepatitis C liver tissue or HCC, antibodies against these proteins, and inhibitors CC fithe expression of groups of genes that are overexpressed in chronic hepatitis C liver tissue or HCC. Groups of genes differentially expressed in chronic hepatitis C liver tissue or HCC may be used for the diagnosis and CC in chronic hepatitis C tissue or HCC may be used for the diagnosis and CC treatment of these diseases. Such genes, inhibitors of their expression CC development of drugs to treat chronic hepatitis C and/or HCC. Sequences CC ABV0459-ABV04790 are SAGE tags representing the 100 least highly corporated with chronic hepatitis C liver tissue.

CC hepatocellular carcinoma compared with chronic hepatitis C liver tissue.
                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
SAGE tag; serial analysis of gene expression; human; Th1 cell;
                                Human Thi cell preferentially expressed EST SAGE tag, SEQ ID NO:170.
                                                                        29-NOV-2002
                                                                                                             ABV78459
                                                                                                                                                                                                                                                                                                                                                                                       Sequence 10 BP; 1 A; 2 C; 6 G; 1 T; 0 other;
                                                                                                                                               ABV78459 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to SAGE (serial analysis of gene expression)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 46;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human chronic hepatitis C tissue expression exasperating comprises 100 high-ranking genes -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-631294/68
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       expression pattern;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SAGE tag; serial analysis of gene expression; human; chronic hepatitis C; CH; liver tissue; hepatocellular carcinoma; cancer; tumour; HCC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human gamma-glutamyltransferase 1 SAGE tag #599
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                                                                                                                                                                                                                                                                  5 GGGCCAGG 12
                                                                                                                                                                                                                                                                                                                Similarity
8; Conserv
                                                                                                                                                                                                                                           GGGCCAGG 10
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                                                                                                                                                                                                                                                                                                                  Conservative
                                                                      (first entry)
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                                                                                                                                               CDNA; 10
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                                                                                                                                                                                                                                                                                                                                   100.0%;
                                                                                                                                                                                                                                                                                                                                                    40.0%;
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                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                               DB 1; Ler
No. 1.1e+02;
                                                                                                                                                                                                                                                                                                                                                    Length 10;
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RESULT 220
ABL57664
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                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 8
                                                                                                                                               gene therapy; respiratory inflammatory disease; polymorphism; primer;
primer extension; ss:
                                                                                                                                                                            SCYA24; human; small inducible cytokine;
                                                                                                                                                                                                        Human SCYA24 primer-extension oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            activated T cell; T lymphocyte; immune response; expression pattern; preferential expression; immune disorder; EST; expressed sequence ta
08-SEP-2000; 2000US-231129P
                                                                                       WO200220851-A1
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                                                                                                                                                                                                                                                                     ABL57664;
                                                                                                                                                                                                                                                                                                 ABL57664 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 10 BP; 4 A; 2 C; 4 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim
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                           10-SEP-2001; 2001WO-US28328
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                                                                                                                   Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      activated Th1 and Th2 cell expression gene group, useful for sais and treatment of Th1 and Th2-related diseases -
                                                                                                                                                                                                                                                                                                                                                                                                  11 GGCCAAAG 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Ler
o. 1.1e+02;
                                                                                                                                                                            isogene; antiasthmatic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 10;
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Solute
KW Gluten
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PR 30-NO\
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AC GENA-
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NOVel
PT Amino
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a novel isolated polynucleotide comprising a small inducible cytokine subfamily A (cys-cys), member 24 (SCYA24) isogene. The polypeptide of the invention has antiasthmatic activity. The polynucleotide may have a use in gene therapy. The polynucleotide and polypeptide are useful in the the development of drugs for treating diseases associated with SCYA24 activity, e.g. respiratory inflammatory diseases such as asthma. The 3' end sequences of oligonucleotide primers used for detecting polymorphisms in the SCYA24 gene by primer extension are represented in ABL57646-ABL57675.
                                                                                                                                                                 Novel genetic variants of Solute Carrier Family 1 (Glutamate/Neutral Amino Acid Transporter), Member 4 isogenes, for improving efficiency and reliability in drug development for treating cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Solute carrier family 1; SLC1A4; haplotyping; human; cancer; priglutamate/neutral amino acid transporter; neurological disease; amino acid transporter disorder; single nucleotide polymorphism
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                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-NOV-2000; 2000US-250254P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-NOV-2001; 2001WO-US44781
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8; Conserv
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                                                                                                              Page
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                                                                                                              16; 139pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Han J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
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                                                                                                                                                                                                                                                                                                                                                  Sausker
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o. 1.1e+02;
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The invention relates to an isolated polynucleotide (I) comprising a first nucleotide sequence which comprises solute carrier family 1 (glutamate/neutral amino acid transporter), member 4 (SLC1A4) isogen

isogenes

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RESULT 222
ABK72637/C
ID ABK726
XX ABK726
XX ABK726
XX Jeukot
XX Leukot
XX Leukot
XX Humman
KW Chemoa
KW inflam
KW Primer
XX Homo s
XX Leukot
XX L
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Best Local S
Matches 8
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                                                                                                                                                                                                                            Novel isolated human leukotriene B4 receptor polynucleotide, therapeutic purposes, for studying expression and function opolynucleotide, and for expressing the receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SICIA4 gene allele specific oligonucleotides, oligonucleotides and related PCR primers used nucleotide polymorphisms (SNP) of the gene.
                                                                                                                                       Claim 17; Page 14; 69pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bieglecki KM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chemoattractant; inflammation; immune response;
inflammatory disorder; recombinant non-human an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABK72637 standard; DNA; 10 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Leukotriene B4 receptor primer extension oligonucleotide #9
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Koshy B,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chemokine receptor-like 1; LTB4R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sanchis A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Ler
o. 1.1e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          animal;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sausker EA;
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The invention describes an isolated (chemokine receptor-like 1) (LTB4R)

human leukotriene B4 receptor polynucleotide (I) comprising

The invention describes an isolated human pyridoxal (pyridoxine, vitamin B6) kinase, (PDXK) polynucleotide. The polynucleotide is useful in studying the expression and function of PDXK, and in expressing PDXK protein for use in screening for candidate drugs to treat PDXK related diseases and for therapeutic purposes. A transgenic animal is useful for studying expression of the PDXK isogenes in vivo, for in vivo screening and testing of drugs targeted against PDXK protein, and for testing the efficacy of therapeutic agents and compounds for autoimmune polyglandular disease type 1. The polypeptide is useful for studying the effect of the variation on the biological activity of PDXK and the binding affinity of

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RESULT 223
ABK17015/c
ID ABK170
AC A
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                                                                                                                                                                                                                                                                                                                                                                                                        Isolated human pyridoxal (pyridoxine, vitamin B6) kinase polyNts useful for therapeutic purposes, for studying the expression and function of the polyNt, and for expressing pyridoxal protein -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pyridoxal kinase; pyridoxine; vitamin B6; PDXK autoimmune polyglandular disease type 1; transgenic gene therapy; primer extension; primer; ss.
                                                                                                                                                                                                                                                                                                                                                 Claim 19; Page 14; 135pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                oligonucleotide used for detecting polymorphisms in the leukotriene B4 receptor (LTB4R) gene.
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8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENAISSANCE PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Duda A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCGGGCCA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (Pyridoxine, vitamin B6) Kinase
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100.0%; Pred. No: 1.1e+02;
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Best Local :
The invention describes an isolated polynucleotide comprising a nucleotide sequence which is a polymorphic variant of a reference sequence for the aldehyde dehydrogenase 5 family, member Al (succinate-semialdehyde dehydrogenase) (ALDH5Al) gene or its fragment. The polymorphic variant with a candidate agent and contacting the ALDH5Al polymorphic variant with a candidate agent and assaying for binding activity. The polymorphide and haplotypes are useful for identifying an association between a trait such as a clinical response to a drug targeting ALDH5Al and a haplotype ALDH5Al gene. Transgenic animals are also useful for studying expression of the ALDH5Al isogenes in vivo, for in vivo screening and testing of drugs against ALDH5Al protein and for testing the efficacy of therapeutic agents and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   polyglandular disease type 1. Genotyping and haplotyping is useful for improving the efficacy and reliability of several steps in the discovery and development of drugs for treating diseases associated with PDXK activity, e.g., autoimmune polyglandular disease type 1, to validate PDXK as a candidate agent for treating a specific condition or disease predicted to be associated with PDXK activity, and in the design of clinical trials of candidate drugs. This sequence is one of 38 (see ABK16978-ABK17015) primers used for detecting PDXK gene polymorphisms by primer extension terminates, described in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                New genetic variants of human aldehyde dehydrogenase 5 family, member Al, ALDH5Al gene for treating metabolic diseases and for expressing ALDH5Al protein useful in identifying drugs to treat 4-hydroxybutyric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antisense technology; primer extension oligonucleotide;
4-hydroxybutyric aciduria; metabolic disease; transgenic animal;
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                                                                                                                                                                                                                                                                                                                                                                               Claim 18; Page 15; 151pp; English
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                     The invention relates to an isolated human superoxide dismutase 1, csoluble (amyotrophic lateral sclerosis 1 (adult)) (SOD1) polynucleotide (I) comprising a sequence which is a polymorphic variant of a reference (C (I) comprising a sequence which is a polymorphic variant of a reference cc sequence for the SOD1 gene. Haplotyping the SOD1 gene of an individual, involves: (a) determining whether the individual has one of the SOD1 (C haplotypes or haplotype pairs given in the specification; or the individual, or the identity of the nucleotide at two or more polymorphic sites selected from PS1-7. The method is useful for determining whether an individual C from PS1-7. The method is useful for determining whether an individual has a haplotype or haplotype pairs defined in the specification. The composition of the discovery and development of drugs for treating consecuted with SOD1 activity, e.g., amyotrophic lateral consecution or disease associated with SOD1 activity. It can for the desire of the security of the condition or disease associated with SOD1 activity. It can for the security of the security of the condition or disease associated with SOD1 activity. It can for the security of the security 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Isolated human superoxide dismutase 1 (SOD1) soluble polynucleotide useful for screening therapeutic compounds, comprises a sequence wh is a polymorphic variant of reference sequence for the SOD1 gene or
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b. 1.1e+02;
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RESULT 226
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XX Human;
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KW Hodgki
KW Iysoso
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Matches 8
                     The invention comprises the human lysosomal acid phosphatase 2 (ACP2) nucleic acid and protein sequences. Specifically, the invention relates to the discovery of 22 novel polymorphic sites within the APC2 gene. The invention also comprises methods for haplocyping and genocyping the ACP2 gene in an individual. The ACP2 gene (located on chromosome 11) encodes a lysosomal-specific enzyme that catalyses the hydrolysis of orthophosphoric monoesters to alcohol and phosphate. The ACP2 gene and protein are pharmaceutically important in the treatment of Hodgkin's disease (HD) and acid phosphatase deficiency. The novel ACP2 gene polymorphisms of the invention are useful in haplotyping the ACP2 gene polymorphisms of the invention are useful in haplotyping the ACP2 gene polymorphisms are disease and acid phosphatase deficiency). The ACP2 gene polymorphisms are useful for ACP2 gene polymorphisms are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   treating a specific condition or disease predicted to be associated with SOD1 activity. (1) is useful in studying the expression and function of SOD1, and in expressing SOD1 protein for use in screening for candidate drugs to treat diseases related to SOD1 activity. AAS95660-AAS95710 represent human superoxide dismutase 1, soluble (amyotrophic lateral sclerosis 1 (adult)) (SOD1) allele-specific oligonucleotides and related PCR primers as described in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel genetic variants of acid phosphatase 2, 1) gene useful in studying expression and function
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; ss; lysosomal acid phosphatase 2; ACP2; gene; chromosome 11; lysosome-specific enzyme; orthophosphoric monoester hydrolysis; lydokin's disease; HD; acid phosphatase deficiency; novel polymorphic site; ACP2 haplotype; ACP2 genotype; polymorphism; transgenic animal; primer; probe; primer-extension oligonucleotide; SNP; single nucleotide polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-JUN-2000;
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8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            etic variants of acid phosphatase 2, lysosomal polypeptide ul in studying expression and function of the protein, and drugs to treat diseases e.g. Hodgkin's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 15; 109pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Messer C,
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                                                                                     The present invention describes a human maturation/activation dendritic cell (DC) expression gene group consisting of 100 genes which show the highest expression among the genes expressed in human maturation/ activation DC. Also described are: (1) a protein expressed by the above human maturation/activation DC expression gene; (2) an antibody against the protein; and (3) an antagonist against the expression of each gene belonging to the above gene group. The gene group is useful for the treatment and the diagnosis of various human diseases related to human DC. ABL42627 to ABL42926 represent specifically claimed human maturation/activation DC expression gene tags from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the invention are useful in the production of a transgenic animal which expresses ACP2 protein. The ACP2 nucleic acids of the invention are useful in the production of allele-specific oligonucleotides designed to genotype each of the ACP2 polymorphisms. Nucleic acids ABL36299-ABL36320 represent claimed ACP2 allele-specific probes. Nucleic acids ABL36231-ABL36364 represent claimed ACP2 allele-specific probes. Nucleic acids ABL36531-ABL36364 represent claimed ACP2 primers. Nucleic acids ABL3654-ABL36408 represent claimed ACP2 primer-extension
                                                            Sequence 10
                                                                                                                                                                                                                                                               Claim 10;
                                                                                                                                                                                                                                                                                            Human maturation/activation dendritic cell expression gene group
                                                                                                                                                                                                                                                                                                                                                                                     22-MAY-2000; 2000JP-0150562
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Similarity
8; Conserv
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                                                                                                                                                                                                                                                           Page 12; 41pp; Japanese
                                                           BP; 3
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.larity 100.0%;
Conservative 0;
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Score 8; DB 1 Pred. No. 1.1 O; Mismatches
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RESULT 229
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                                                                                           24-JUL-2002.
                                                                                                                                                                                                                                                              Human; Kaposi's
                                                                                                                                                                                                                                                                                                              Kaposi's Sarcoma SAGE library Tag No.15 SEQ ID NO:15.
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                             23-JAN-2002; 2002EP-0075264
                                                                                                                                                                                                      Homo sapiens
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes a method for determining if an individual has a tumour cell or site of angiogenesis, or if a treatment is effective in changing angiogenesis or changing a status of a set of target cells, comprising determining if a sample of the subject has an expression product of at least one marker gene. Also described is a compound capable of altering the expression or activity of Keratin 14, TIE 1, Salioadhesin or Siglec in a cell. Peripheral blood mononuclear cell (PHMC) expressed Keratin 14, TIE 1, Salioadhesin or Siglec, and kits containing them from the present invention can be used in a diagnostic method, particularly as an indicator of angiogenesis or to determine presence of a tumour cell. The method of the invention is suitable to determine within a few days if a certain treatment against Kaposi's Sarcoma is successful. AB081851 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                           WPI; 2002-590638/63
                                                                                                                    Petersohn D,
                                                                                                                                                                                                                                            03-JAN-2001; 2001DE-1000127
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28-SEP-2001; 2001EP-0203703
28-SEP-2001; 2001US-325722P
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Pred. No. 1.4e+02;
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In vitro identification of skin-expressed genes, useful for determining

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RESULT 231
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                  The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin
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disorders,
                                                                                                                                                                                                                                                               Disclosure; Page 217; 1345pp;
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      neurodermatitis;
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ving cosmetic or pharmaceutical agent
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Pred. No. 1.4e+02;
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                                                                                                                                                                                                                                                                                                                                             The invention relates to identifying (M1) genes in vitro that, in humans or animals, are important for skin ageing and/or skin stress by serial analysis of gene expression between mixtures of transcribed and optionally translated, genetically encoded factors (A) obtained from young and aged skin, to identify that genes that show strong differential expression. (A) comprises protein or mRNAs or their fragments. (M1) is
                                                                                                                                                                                                           expression. (A) comprises protein or mRNAs or their fragments. (M1) is useful for: identifying markers of skin ageing and/or stress; determining skin ageing and/or stress; and identifying or determining the effects of pharmaceutical or cosmetic agents for control of skin ageing. The present sequence is one of a group of human skin ageing/stress related expressed sequence tags (ABQ86246-ABQ87680) of the invention.
                                                                                                                                                                  Sequence 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying genes involved in skin stress and ageing, useful e.g. screening for cosmetic or therapeutic agents, based on differentia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Petersohn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-JAN-2001; 2001DE-1000121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; skin ageing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human skin stress/ageing related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-SEP-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma skin. The present sequence is that of a human expressed sequence (EST) of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus;
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                                                                                      Similarity
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     cceeeccaeec 13
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                                                                                                                                                               BP; 0 A; 4 C; 5 G; 2 T; 0 other;
                                                           Conservative
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81.8%;
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                                                                                      Score 7.8; DB 1;
Pred. No. 1.4e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred.
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                                                      Mismatches
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                                                                                                         Length 11,
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                                                           Gaps
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RESULT 233

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ABZ72939/c
ID ABZ729
                                                                                     The present invention describes a recombinant adeno-associated virus CC (AAV) vectored ribozyme composition (1) (1) comprises; (a) at least a CF first ribozyme that specifically cleaves an mRNA encoding a protein, CC polypeptide, or peptide selected from the group of rod opsin, iNOS, CR RDS/peripherin, VEGFR1, VEGFR2, adenosine A-2B receptor, IGF-1, integrin alpha 1, integrin alpha 3, integrin alpha 5, or integrin alpha 10; (b) a CC vector comprising a polynucleotide encoding the ribozyme. Where the polynucleotide operably positioned downstream of at least a first CC promoter that directs expression of the polynucleotide in a selected CC mammalian cell transformed with the vector; (c) a viral particle comprising the ribozyme or the polynucleotide; (d) an AAV vector CC comprising the ribozyme or the polynucleotide; or (e) a host cell CC comprising the ribozyme or the polynucleotide. Also described is a method CC for decreasing the amount of mRNA encoding a selected polypeptide in a CC retinal cell of a mammalian eye, comprising to the eye the CC composition described above, and for a time effective to specifically cleave the mRNA in the cell. (I) has ophthalmological activity, and can be used in gene therapy. (I) can be used for treating a disease or CC dysfunction of the mammalian eye, such as a retinal disease or treinal disease or treinal disease or retinal disease or retinal dominant for treating the diseases mentioned above, including autosomal dominant retinitis or a blood-retinal barrier dysfunction. (I) can also be useful CC pathological condition, e.g. atrophic or pigmented lesions of the eye blood of the mreach invention or pigmented lesions of the eye.

CC constitution of the mreach invention or pigmented lesions of the eye.
  Query Match 39.
Best Local Similarity 81.
Matches 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A recombinant adeno-associated virus-vectored ribozyme composition, useful for treating a disease or dysfunction of the mammalian eye e.g. retinal disease, e.g. diabetic retinopathy or age-related macular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hairpin ribozyme; hammerhead ribozyme; ribozyme; retinal disease; target; ophthalmological; gene therapy; eye; retinal dysfunction; AAV; diabetic retinopathy; macular degeneration; autosomal dominant retinitis; blood-retinal barrier dysfunction; adeno-associated virus; blindness; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-MAY-2001; 2001US-0847601.
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                                                                                                                       exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 5; Page 73; 115pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-111880/10
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                                                                                 A; 4 C; 4 G; 1 U; 0 other;
                 39.0%;
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Score 7.8; DB 1;
Pred. No. 1.7e+02;
0; Mismatches 2
                                      Length 12;
  Indels
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Gaps
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RESULT 234
AADD3853/C
ID AADD3853/C
ID AADD3853/C
ID AADD38
XX ADD38
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XX ADD38
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XX Cell 9
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XX US6218
XX US6218
XX O5-SEP
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RESULT 235
AAS18424/C
ID AAS184
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DT 12-MAR
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                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is a degenerate PCR primer used for amplifying the human cell cycle checkpoint protein, hohkl DNA. The cell cycle checkpoints are regulatory pathways that control the order and timing of cell cycle transitions, and ensure that critical events such as DNA replication and chromosome segregation are completed with high fidelity. The chkl protein controls cell cycle in response to DNA damage. It functions as kinase and phosphorylates the key regulators of Cdk tyrosine phosphorylation. The checkpoint gene sequences are used as probes for a portion of the chromosome associated with tumours and other malignancies, as well as growth and/or development deficiencies. The chkl proteins are useful for generating specific antibodies and for inhibiting growth of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; cell cycle checkpoint; chk1; tumour; malignancy; cell growth inhibitor; development deficiency; PCR primer; DNA damage; kinase; ss.
                                                                                                             AAS18424 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New Chk1 proteins and gene sequences encoding the proteins use probes for a portion of the chromosome associated with tumors malignancies, growth and/or development deficiencies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 17
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9; Conserve
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                                                                                                                                                                                                                                                               GCCTGTCCAA 1
                                                                                                                                                                                                                                                                                                      GGCCAGGCCAA 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Column 24; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 5 A; 4 C;
                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sanchez
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97US-0924183.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97US-0924183
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                                                                                                                                                                                                                                                                                                                                                                                                          39.0%;
81.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                          Score 7.8; DB 1;
Pred. No. 2.4e+02;
                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 17;
                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            useful
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                                                                                                                                                                                                                                                                                                                                                                               Gaps
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other
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12-MAR-2002 AAS18424;

(first entry)

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RESULT 236
ABX16354/c
ID ABX16354;
XX
AC ABX16354;
XY
DT 08-APR-2003 (first of the control of th
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Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         checkpoint (chk1) proteins and the nucleic acid sequences encoding them. Human chk1 (hchk1) maps to chromosome 11q24. Chk1 is involved in cellular responses to DNA damage, in the cell cycle checkpoint pathway. The protein is useful for generating specific antibodies and for inhibiting the growth of cells. The nucleotide sequence encoding the protein may be used as a probe for a portion of the chromosome associated with tumours and other malignancies, as well as growth and/or development deficiencies. The present sequence represents a degenerate
                                  12-DEC-2001; 2001US-0020038
                                                                                                                                                                                                                                                                                       Human; checkpoint; chk1; anti-Chk1 antibody;
                                                                                                                                                                                                                                                                                                                                              Human checkpoint gene Chk1 PCR primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 5 A; 4 C; 5 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New mammalian checkpoint protein and gene, for generating specific antibodies or for inhibiting the growth of cells, and for use as a probe for a portion of a chromosome associated with tumours or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; chec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to the isolation of human and mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Column 24; 39pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-JAN-2000; 2000US-0488364
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primer used to amplify DNA encoding human chk1 protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (BAYU ) BAYLOR COLLEGE MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity nes 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2002-040207/05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GGCCAGGCCAA 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GCCTCTCCAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  checkpoint pathway; inhibition of cell growth; tumour; growth deficiency; development deficiency; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                        (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97US-0924183
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               39.0%;
81.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 7.8; DB 1;
Pred. No. 2.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amplify DNA encoding human chk1
                                                                                                                                                                                                                                                                                          tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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RESULT 237
AAX293
AXX AAX293
XX AAX293
XX AAX1
DT 10-JUN
XX AAX1
KW ANTISE
KW ANTISE
KW HYPETP
OS SYNTHE
OS HOMO S
XX WO9909
XX SYNTHE
OS HOMO S
XX I SIS-
FEB
PR 07-AUG
PR 13-AUG
XX 11SIS-
PA (ISIS-
PA WPI; 1
XX AAX1
XX WPI; 1
XX TSIS-
XX WPI; 1
XX AAX1
XX WPI; 1
XX WPI; 1
XX TSIS-
XX WPI; 1
XX 
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Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention describes an anti-Chkl antibody capable of specifically binding to an antigenic determinant on the proteins encoded by a sequence comprising 476 (3 sequences), 479, 496 or 513 amino acids. A new method is used to produce the antibody, which is useful for detecting a Chkl protein that is associated with a tumour. This sequence represents a PCR primer used to isolate DNA encoding the
                                                      Example 4; Page 92; 190pp;
                                                                                                       New antisense oligonuclectides that detect and modulate expression of Jun N-terminal kinase proteins - useful f hyperproliferative diseases and inhibiting tumor growth and for modulating protein phosphorylation by these protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 13; 28pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New anti-Chkl antibody, that may be a monoclonal or polyclonal antibody, useful for detecting a Chkl protein that is associat
                                                                                                                                                                                                                                            WPI; 1999-181060/15
                                                                                                                                                                                                                                                                                                                                                   (ISIS-)
                                                                                                                                                                                                                                                                                                                                                                                                    13-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9909214-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antisense oligonucleotide; Jun N-terminal kinase; JNK; hybridise; JNK2; JNK3; cell cycle progression; phosphorvlation: tumour: proba
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chemically modified sense control
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAX29342 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human checkpoint protein Chk1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-182651/18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-JAN-2000; 2000US-0488364.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11 GGCCTGTCCAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          l Similarity
9; Conserva
                                                                                                                                                                                                                                                                                                                                                   ISIS PHARM INC.
                                                                                                                                                                                                                                                                                             Gaarde WA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 5 A; 4 C; 5 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sanchez
                                                                                                                                                                                                                                                                                                                                                                                                    97US-0910629
                                                                                                                                                                                                                                                                                                                                                                                                                                                        98WO-US16488
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cycle progression; phosphorylation; tumour; probe;
/e disease; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entry)
                                                                                                                                                                                                                                                                                             Mckay R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     39.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
                                                      English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ВР
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Pred. No. 2.4e+02;
                                                                                                                                                                                                                                                                                             Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              probe ISIS No:
                                                                                                                                                                                                                                                                                                Nero
                                                                                                         tumor growth in as
by these proteins

    useful for treating
mor growth in animals,

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2
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                                                                                                                                                          for treating
                                                                                                                                                                                           the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JNK1;
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The invention relates to antisense oligonucleotides that detect and

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RESULT 238
AAX29331/C
ID AAX293
AC AAX293
AC AAX293
AC AAX293
XX IO-JUN
DT 10-JUN
DT 10-JUN
DX JNK2-8
XX Antise
XW JNK2;
XW hyperp
XX Synthe
OS Homo s
XX WO9909
PN WO9909
XX 25-FEB
PD 13-AUG
XX (ISIS-
PR 13-AUG
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Best Local S
Matches 10
modulate the expression of Jun N-terminal kinase (JNK) proteins. The oligonucleotides specifically hybridize to a nucleic acid encoding a JNK1, JNK2 or JNK3 protein, and which modulate expression of these proteins. The oligonucleotides are useful for modulating JNK protein expression and cell cycle progression in cultured cells or animal cells. The oligonucleotides are also useful for modulating the phosphorylation of a protein that has been phosphorylated by a JNK protein, and the expression of a cellular protein that promotes one or more metastatic events. The oligonucleotides also form pharmaceutical compositions for treating animals with a hyperproliferative disease, and for inhihiming
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modulate the expression of Jun N-terminal kinase (JNK) proteins. The oligonucleotides specifically hybridize to a nucleic acid encoding a JNK1, JNK2 or JNK3 protein, and which modulate expression of these proteins. The oligonucleotides are useful for modulating JMK protein expression and cell cycle progression in cultured cells or animal cells. The oligonucleotides are also useful for modulating the phosphorylation of a protein that has been phosphorylated by a JMK protein, and the expression of a cellular protein that promotes one or more metastatic events. The oligonucleotides also form pharmaceutical compositions for treating animals with a hyperproliferative disease, and for inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense oligonuclectides that detect an expression of Jun N-terminal kinase proteins hyperproliferative diseases and inhibiting tu and for modulating protein phosphorylation by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-181060/15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9909214-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense oligonucleotide; Jun N-terminal kinase; JNK; hybridise; JNK1; JNK2; JNK3; cell cycle progression; phosphorylation; tumour; probe; hyperproliferative disease; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JNK2-specific probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-JUN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAX29331 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                  Example 4; Page 87; 190pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-AUG-1998;
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Pred. No. 2.3e+02;
0; Mismatches 4
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Best Local
                                                                                                                                                                                                                                                 induction of apoptosis. The present sequence is one such antisense oligonucleotide. The oligonucleotides of the present invention are useful for treating diseases or conditions with reduced apoptosis, e.g. cancer and cellular hyperproliferation. The oligonucleotides may also be used to increase the stimulation of apoptotic proteins, e.g. for treating Alzheimer's or Parkinson's disease, amylotrophic lateral sclerosis, retinitis, pigmentosa, epilepsy, myocardial infarction, stroke, obstructive jaundice, polycystic kidney and diabetes. The present sequence may have a phosphorothicate backbone.
                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to antisense oligonuclectides (AAC62844-C63000, AAA96093-A96099 and AAA07993) that hybridise specifically to a nucleotide encoding a Jun N-terminal kinase (JN protein, resulting in decrease of JNK2 expression and leading to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense; gene therapy; JNK2 protein; apoptosis; cancer; cellular hyperproliferation; Alzheimer's; Parkinson's disease; amylotrophic lateral sclerosis; retinitis; pigmentosa; epilepsy; myocardial infarction; stroke; obstructive jaundice; polycystic kidney;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel methods for reducing antisense oligonucleotides,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diabetes; Jun N-terminal kinase; ss.
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                                                GTCCGGGCCAGGCC
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GGCCTGGCCCGGAC
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71.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  apoptosis comprising contacting cells with useful for treating apoptotic disorders,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВP
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                                                                                                            Score 7.6; Ub .,
No. 2.3e+02;
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Pred. No. 2.3e+02;
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AC AAA486
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DT 20-SEP
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AAC62885
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antisense; gene therapy; JNK2 protein; apoptosis; cancer; cellular hyperproliferation, Alzheimer's; Parkinson's disease; amylotrophic lateral sclerosis; retinitis; pigmentosa; epilepsy; myocardial infarction; stroke; obstructive jaundice; polycystic kidney; diabetes; Jun N-terminal kinase; ss.
              20-SEP-2000
                                          AAA48651;
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                                                                                                                                                                                                                                                       Sequence 20 BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                             specifically to a nucleotide encoding a Jun N-terminal kinase (J) protein, resulting in decrease of JNK2 expression and leading to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               e.g. cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel methods for reducing apoptosis comprising antisense oligonucleotides, useful for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to antisense oligonucleotides (AAC62844-C63000, AAA96093-A96099 and AAA07993) that hybridise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 4; Page 135; 160pp; English.
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                                                                                                                                                                                                                                                                                   sequence may have a phosphorothioate backbone.
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Pred. No. 2.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            contacting cells with apoptotic disorders,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WA;
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                                                                                                                                                                                                                                                                                                                                                                                                                              (JNK2)
                                                                                                                                                                                                Gaps
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present sequence is the JNK2 antisense oligonucleotide. The antisense oligonucleotides used in the method contained modifications, namely phosphorothioate linkages and 2 methoxyethoxy bases. Some C residues also had a 5 methyl modification. Inhibitors of the TNF alpha signalling molecules have antibacterial, immunosuppressive, antipsoriatic, antidiabetic, antithyroid, cytostatic, dermatological, antiallergic and antiinflammatory activity. The antigense inhibitors may be useful for the treatment of sepsis, rheumatoid arthritis, inflammatory, immune disease, inflammatory bowel disease, allergic
                                                                                                                                                                                                                        A novel method for modulating cell adhesion molecule expression involves antisense inhibition of a tumour necrosis factor (TNF) alpha signalling molecule. In the method TNF alpha signalling molecules Ha-ras, c-raf and c-Jun N-terminal kinase (JNK) 2 were inhibited by antisense oligonucleotides. In addition an antisense oligonucleotide to the cell adhesion molecule E-selectin was also examined. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antisense; E-selectin; TNF alpha; cell adhesion; tumour necrosis factor alpha; phosphorothioate, methoxyethoxy; sepsis; rheumatord archritis; inflammatory; immune disease; inflammatory bowel disease; allergic contact dermatitis; psoriasis; diabetes; Grave's disease; allograft rejection; cancer; antibacterial; immunosuppressive; antipacriatic; antidiabetic; antithyroid; cytostatic; dermatological; antiallergic; Ha-ras; c-raf;
                                                                                                                                                                                                                                                                                                                                                                              Modulating cell adhesion molecule expression for treating immune or inflammatory diseases involves treating cell with specific inhibitor Tumour Necrosis Factor alpha signalling molecule
                                                 contact dermatitis, psoriasis, diabetes, Grave's
                                                                                                                                                                                                                                                                                                                                                 Claim 36; Page 46; 100pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200034303-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Кeу
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Antisense oligonucleotide ISIS no.15354 to human JNK2 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000-423367/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N-terminal
                                  and cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              98US-0209668.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99WO-US28965
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /mod_base= OTHER
/note= "All bases are 2'-methoxyethoxy,
additionally C bases are m5c"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/note= "All bases are 2'-methoxyethoxy
additionally C bases are m5c"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "Phosphorothioate internucleotide linkage"
16..20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mod
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JNK;
                                                  disease, allergic disease, allograf
                                                                                                                                                                                                                                                                                                alpha
                                                                                                                                                                                                                                                                                                                                                                                                      of.
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Query Match
Best Local Similarity
Matches 10; Conserv

Conservative

<u>,</u>

38.0%; 71.4%;

Score 7.6; Pred. No. 2 Mismatches

DB 1; 2.3e+02;

Length

Indels

0

Gaps

0

Sequence

20

BP; 4 A;

7 C; 7 G; 2 T; 0 other;

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                                                                                                                            Query Match
Best Local S
Matches 10
                                                                                                                                                                                              The present invention relates to the use of coligonucleotides for treating cancer and for mimic or augment the effect of JNK antisense mimic cancer. The present sequence is one such
                                                                                                                                                                                                                                                                 Use of Jun Kinase antisense mRNA for treating cancer by administering vector comprising promoter operably linked to DNA sequence that encodes the antisense mRNA to patient diagnosed with cancer \,\cdot\,
 12-JUN-2001
                   AAF99183;
                                      AAF99183 standard;
                                                                                                                                                                    Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JNK; jun kinase; antisense; cytostatic;
2'-O-methoxyethyl oligonucleotide; MOE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAH23754;
                                                                                                                                                                                       oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                           10-NOV-2000; 2000WO-US30869
                                                                                                                                                                                                                                                                                                                                                                                                              17-MAY-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200134792-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JNK1 antisense oligonucleotide, JNK2AS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAH23754
                                                                                                                                                                                                                                                                                                                                                   (USSH ) US
                                                                                                                                                                                                                                                                                                                                                                       12-NOV-1999;
                                                                                                                                                                                                                                                                                                             2001-335925/35
                                                                                       14 GGCCTGGCCCGGAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14
                                                                                                         10;
                                                                                                                                        Similarity
                                                                                                         GTCCGGGCCAGGCC 14
                                                                                                                                                                                                                                                  Page 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GTCCGGGCCAGGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GGCCTGGCCCGGAC
                                                                                                                                                                                                                                                                                                                                                   DEPT HEALTH & HUMAN SERVICES
                                                                                                                                                                    BP; 4 A;
                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                              Gorospe
 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                                                                      99US-0165224.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  //note= "This oligonucleotide is a 2'-O-methoxyethyl
chimeric antisense oligonucleotide containing five
MOB/phoshopdiester residues flanking a
2'-deoxynucleotide/phosphorothioate region"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                75pp; English.
                                      DNA;
                                                                                                                                      38.0%;
71.4%;
                                                                                                                                                                    7 C; 7
                                                                                                                                                                                                                                                                                                                                3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           _base= "OTHER"
                                      20
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                                                                                                                                                                                                                                                                                                                                 Holbrook NJ;
                                       ВÞ
                                                                                                                                                                    <u>ن</u>
                                                                                                                            Score 7.6; DB
Pred. No. 2.3e
0; Mismatches
                                                                                                                            0
                                                                                                                                                                    N
                                                                                                                                                                    T; 0
                                                                                                                                                                    other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ISIS #12560)
                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               phosphorothioate;
                                                                                                                                                                                                         of Jun Kinase (JNK) antisense for screening compounds that ense oligonucleotides treatmen
                                                                                                                                                                                                 JNK antisense
                                                                                                                                               Length 20;
                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                88
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                                                                                                                            Gaps
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RESULT 244
ABS77827/c
ID ABS778
XX
AC ABS778
XX
AC ABS778
XX
DT 13-DEC
XX
DE Angiog
XX
Angiog
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                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 10
             Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth, tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy, retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular gla
                                                                                                                                                                                                                                                                                                                                                                                                                                                      against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), antigens (e.g. toxoplasma, and/or paramaly, campylobacter, clostridium, Escherichia coli and/or tampylococcus), fungal antigens and/or parasitic antigens. The method is staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an
                                                                                             Angiogenesis inhibitory oligonucleotide #311.
                                                                                                                               13-DEC-2002
                                                                                                                                                                                             ABS77827
                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immune response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py_rich) or thymidine (T) rich. The method is used to vaccinate subjects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 101; Page 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vaccinating against tumors, infectious diseases, allergies using immunostimulatory Py-rich and TG nucleic acids -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Krieg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-SEP-1999;
27-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   immunostimulatory; tumour; viral infection; bacterial infection;
fungal infection; parasitic infection; cancer; asthma;
infectious disease; allergy; immune deficiency; phosphorothicate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-AUG-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Vaccine; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunostimulatory nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (IOWA ) UNIV IOWA RES FOUND (COLE-) COLEY PHARM GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                            the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AΜ,
                                                                                                                                                                                                                                                                             14.
                                                                                                                                                                                                                                                                                                           1 GTCCGGGCCAGGCC 14
                                                                                                                                                                                                                                                                                                                                            10;
                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                             GGCCTGGCCCGGAC 1
                                                                                                                                                                                                                                                                                                                                                                                                            20
                                                                                                                                                                                             standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                         present sequence may have a phosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Schetter C,
                                                                                                                                                                                                                                                                                                                                                                                                         BP; 4 A; 7 C; 7
                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000US-0227436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      99US-0156113.
99US-0156135.
                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      338pp;
                                                                                                                                                                                                                                                                                                                                                         38.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      virucidal;
                                                                                                                                                                                             20
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                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         English
                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                           Score 7.6; I
                                                                                                                                                                                                                                                                                                                                                                                                          G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      #299
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    bactericidal; fungicidal;
                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                            DB
                                                                                                                                                                                                                                                                                                                                                         3e+02;
                                                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                            20
                                                            growth;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      anti-parasitic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and
               glaucoma;
                                                                                                                                                                                                                                                                                                                                            <u>.</u>.
                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        88.
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RESULT 245
ABL39057/c
ID ABL390
XX
AC ABL390
XX Inmuno
DE Immuno
XX
Immuno
XX
Immuno
XX
Antibo
KW Antibo
KW Angiog
XX
Synthe
XX
FT Key
FT modifi
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W02001
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic acid of the invention.
 WO200197843-A2
                                                                                                                                   Synthetic
                                                                                                                                                              Antibody-induced cell lysis; cancer; immunostimulatory; CD20; angiogenesis; metastasis; cytostatic; phosphorothioate backbo
                                                                                                                                                                                                                 Immunostimulatory nucleic acid SEQ ID NO: 463.
                                                                                                                                                                                                                                                    16-APR-2002
                                                                                                                                                                                                                                                                                    ABL39057;
                                                                                                                                                                                                                                                                                                                   ABL39057 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Inhibiting angiogenesis in a subject, involves administering one antiangiogenic nucleic acid molecule to the subject -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis; scleroderma; hypertrophic sca
                                                                              modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-566690/60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-DEC-2001; 2001WO-US48458.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to inhibiting angiogenesis in a subject,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-DEC-2000; 2000US-25534P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                                                                                                                                                                                                                                                                                                                                                                                   14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COLEY PHARM GROUP INC
                                                                                                                                                                                                                                                                                                                                                                                                                                  GTCCGGGCCAGGCC 14
                                                                                                                                                                                                                                                                                                                                                                                                   GGCCTGGCCCGGAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 4 A; 7 C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                (first entry)
                           /*tag= a
/mod_base= OTHER
/note= "phosphorothioate backbone"
                                                                                   1..20
                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           276pp;
                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                38.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            atherosclerosis; scleroderma; hypertrophic scar.
                                                                                                                                                                                                                                                                                                                   20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             English.
                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 7.6; DB 1;
Pred. No. 2.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. 2.30
); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                  backbone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                  88
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RESULT 246
AAX54706/c
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Best Local S
Matches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of cell surface antigens and antibodies. The methods are useful for treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, breast cancer, cervical cancer, colon and rectum cancer, connective tissue cancer, oesophageal cancer, eye cancer, kidney cancer, larynx cancer, leukaemia, liver cancer, lung cancer, Hodgkin's lymphoma, non-Hodgkin's lymphoma, melanoma, oral cavity cancer, ovaricancer, pancreatic cancer, prostate cancer, rhabdomyosarcoma, skin cancer, stomach cancer, testicular cancer, and uterine cancer. The
                                                                                                                            chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer; hepatocellular carcinoma; kidney cancer; melanoma; hepatic metastasis
                                                                                                                                                                                                                               impaired respiration; inflammation; lung disease;
pulmonary vasoconstriction; inflammation; allergic rhinitis;
                                                                                                                                                                                                                                                                                                   Human fibronectin antisense oligonucleotide fragment
                                                                                                                                                                                                                                                                                                                                    05-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to methods for treating or preventing cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression o cell surface antigens and antibodies to a subject having or at risk of
                                                                                                                prostate cancer;
                                                                                                                                                                                pulmonary hypertension; pulmonary vasoconstriction;
                                                                                                                                                                                             acute asthma, allergy; asthma; impeded respiration respiratory distress syndrome; pain; cystic fibros
                                                                                                                                                                                                                                                                  Antisense oligonucleotide; multiple target; antisense treatment;
                                                                                                                                                                                                                                                                                                                                                                                                      AAX54706 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       present sequence is an immunostimulat
the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 212; 312pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         developing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Weiner G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-JUN-2000; 2000US-213346P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-JUN-2001; 2001WO-US20154.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (IOWA ) UNIV IOWA RES FOUND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GTCCGGGCCAGGCC 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GGCCTGGCCCGGAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 4 A; 7 C; 7 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                SS
                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                    entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     38.0%;
71.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunostimulatory oligonucleotide described
                                                                                                                                                                                                                                                                                                                                                                                                        ø
                                                                                                                                                                                                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 7.6; DB 1;
Pred. No. 2.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                  fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                               emphysema
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
                                                                                                                                  metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ovarian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      larynx
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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0

25-MAR-1999

WO9913886-A1

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RESULT 247
AAF20275/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            directed against at least 2 mRNAs selected from target genes, coding and CC non-coding regions of RNAs corresponding to target genes, gene initiation codons, genomic flanking regions, intron-exon borders, the CC initiation codons, genomic flanking regions, intron-exon borders, the CC stand the juxta-section between coding and non-coding CC regions and all segments of RNAs encoding proteins associated with one CC or more diseases, conditions or mixtures. The antisense oligonucleotides CC may be derived from sequences AAX55272-74. These multiple target CC antisense treatment of diseases and conditions. Typical diseases and conditions are those associated with impaired regiration and CC inflammation, including lung diseases, pulmonary vasconstriction, including lung diseases, pulmonary vasconstriction, crepiration, respiratory disease stima, allergies, asthma, impeded CC pulmonary hypertension, pulmonary vasconstriction, emphysema, chronic Obstructive pulmonary disease (COPD), and cancers such as leukemias, CC lymphomas, carcinomas e.g. colon cancer, breast cancer, lung cancer, correlated cancer, hepatocellular carcinoma, kidney cancer, melanoma, for the lungs metastasize and types of cancers and more and types of cance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 8
                                                                                                                                         Low adenosine antisense oligonucleotide; phosphorothioate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic; respiratory obstruction; pulmonary obstruction; impeded respiration; surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pulmonary vasoconstriction; allergic rhinitis; pulmonary hypertension; emphysema; pulmonary transplantation rejection; chronic obstructive pulmonary disease; pulmonary infection; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                    Human fibronectin polynucleotide fragment #1842.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF20275;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 9 BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The specification describes antisense oligonucleotides (AAX52869-X55271) directed against at least 2 mRNAs selected from target genes, coding and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 55; 120pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense oligonucleotides used in treatment of, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1999-229400/19
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17-SEP-1997;
                       WO200062736-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF20275 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               have metastasized to the lungs, including breast and prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
8; Conserv
                                                                                                                         88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        98US-0093972.
97US-0059160.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4 C; 5 G; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   37.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 7.4; DB 1;
Pred. No. 5.2e+02;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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CC oligonucleotides and compositions (I) comprising them. In the antisense (CC oligonucleotides the A is replaced by a 'Universal' or alternative base. (CI) can have respiratory, bronchodilator, antiinflammatory, analgesic, immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. (CC immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. (CC expression and or activity of target polypeptides associated with CC lung/respiratory disorders and malignancies, such as stimulating and cativity of target polypeptides associated with CC lung/respiratory disorders and malignancies, such as stimulating and cativity and antibodies, antibody receptors, cytokines and cativities and indunoglobulins and antibodies, antibody receptors, cytokines and chemokines, endogenously produced specific and non-specific enzymes, chemokines, endogenously produced specific and non-specific enzymes, chemokines, endogenously produced specific and non-specific enzymes, chemokine receptors, adenosine receptors, bradykinin receptors, central conceptors, (CNS and peripheral nervous and non-nervous system peptide creeptors, binding proteins, growth factors, vasoactive peptides and creeptors, binding proteins and malignancy associated proteins. The cantisense oligonucleotides may be used in this way to treat disorders cincluding respiratory obstruction (especially pulmonary obstruction cand/or surfactant hypoproduction which are associated with a disease or condition selected from pulmonary vasoconstriction, inflammation, pathory, pulmonary cathorises, pathory disease or conditions, emphysema, chronic obstructive pulmonary disease (CDD), emphysema, chronic obstructive pulmonary disease (CDD),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Low adenosine (A) content antisense oligonucleotides which trigger adenosine receptors during metabolism, useful e.g. cancers and respiratory obstructions -
                                                                        pulmonary transplantation rejection, pulmonary infections, bronchitis, and/or cancer. AAF18434 to AAF21543 represent human polynucleotide fragments and antisense oligonucleotides used in the exemplification of
                                                                                                                                                         hypertension, emphysema, chronic obstructive pulmonary disease (COPD)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 14; Page 220; 1592pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-MAR-2000; 2000WO-US08020
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                                                  present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EAST CAROLINA
J W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99US-0127958
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              describes low adenosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (A) content antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              g
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Sequence 9 BP; 0 A; 4 C; 5 G; 0 U; 0 other;

Query Match Best Local ( Matches φ 1 GTCCGGGCC 9 Similarity 8; Conserv eccceecc 1 Conservative 88.9%; 37.0%; <u>.</u> Score 7.4; DB 1; Pred. No. 5.2e+02 Mismatches 2e+02; Length Indels 0 Gaps

0

RESULT 248
AAA34153/c
ID AAA341
XX AAA341
XX AAA341
XX AAA341
XX BUMAN
DT 28-JUL
XX HUMAN
XX HUMAN
XX HUMAN
XX HUMAN
XX HUMAN
XX HUMAN
XX A11erg AAA34153 standard; DNA;

ø ВÞ

밁 S

Human adenosine receptor related polynucleotide SEQ ID NO:1842 28-JUL-2000 (first entry)

allergic disease; bronchoconstriction; inhibitor; Human; adenosine receptor; low adenosine antisense oligonucleotide; phosphorothioate; impaired respiration; inflammation; allergy; antiinflammatory;

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RESULT 249
ABQ71850
ID ABQ718
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                                                                                                                                                                                                                                         Query Match
Best Local Similarity
"" hes 8; Conserv?
                                                                                                                                                                                                                                                                                                                                          e.g. ischaemic conditions, pulmonary vasoconstriction, allergies, asthma, impeded respiration, respiratory discress syndrome, pain, cystic fibrosis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPP), and cancers such as leukaemias, lymphomas, carcinomas, and cancers which may metastasise to the lungs, including breast and prostate cancer. The reduction of the adenosine content of the ONs reduces side effects. The A-containing ONs break down with the release of deoxyadenosine which activates adenosine receptors causing bronchoconstriction and inflammation. AAA33313 to AAA33512 represent the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a new composition comprising an antisense oligonucleotide (ON) with low adenosine (up to 15%), which targets nucleic acids involved in bronchoconstriction, allergies, and/or inflammation. The ON can have antiinflammatory, antiallergic, antiasthmatic, cytostatic and analgesic activities. The compositions are useful for the treatment of diseases associated with inflammation, impaired airways, including lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating
  28-AUG-2002
                                         ABQ71850;
                                                                            ABQ71850 standard; DNA; 9 BP
                                                                                                                                                                                                                                                                                                                                          Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                             nucleotide sequences given in the sequence listing from the present invention, which correspond to SEQ ID NO:1 to 2815, and then the last 185 sequences are also called SEQ ID NO:1 to 185, but the sequences differ from the previously named sequences. SEQ ID NO:11 to 1680 (AAA33923 to AAA3392) are specifically claimed ONs from the present invention. N.B. Sequences given in the disclosure of the present invention do not match up with their corresponding SEQ ID NO: sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstriction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; lung disease; ischaemic condition; pulmonary vasoconstriction; asthma; respiratory distress syndrome; pain; cytetic fibrosis; emphysema; pulmonary hypercension; chronic obstructive pulmonary disease; COPD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 495; 1343pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-205971/18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-AUG-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer; leukaemia; lymphoma; carcinoma; metastasis; ss.
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                                                                                                                                                                                                                                                                                                                                                                               in the sequence listing.
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                                                                                                                                                                                                                    ercceecc 9
                                                                                                                                                                                                                                                                                                                                          BP;
                                                                                                                                                                                                                                                             Conservative
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(first entry)
                                                                                                                                                                                                                                                                                                                                      0 A;
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                                                                                                                                                                                                                                                                                                                                          4 C;
                                                                                                                                                                                                                                                                              37.0%;
88.9%;
                                                                                                                                                                                                                                                                                                                                        5 G; 0 U;
                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                         Score 7.4; DB 1;
Pred. No. 5.2e+02;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                          0 other;
                                                                                                                                                                                                                                                                                            Length 9;
                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                           0,
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ABQ71851
ID ABQ
XX
AC ABQ
XX
DT 28-,
XX
DE Zin
XX
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                                                                                                                                                                                                                                                                                                                                              RESULT 250
                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (3) designing (M) (I) involves selecting the F1 zinc finger such that it binds to the S1 target subsite, selecting the F2 zinc finger such that it binds to the S2 target subsite, and selecting the F3 zinc finger such that it binds to the S3 target subsite, thus designing (I) that binds to a target site. (I) is useful for recognition of triplet target subsites having the nucleotide G in the 5 most position of the subsite. (I) is useful in studying gene function, and for human therapeutics and plant engineering. (I), (II) or (III) is useful in therapeutic methods to modulate the expression of a target region within
                                                                                                                    ABQ71851 standard; DNA; 9 BP
                                                                                                                                                                                                                                                                                                                                                                                          a subject, in diagnostic methods for sequence specific detection of target nucleic acid in a sample, and in assays to determined the phenotype and function of gene expression. (I) has improved affinity and specificity for their target sequences, as well as enhanced biological activity. ABQ71213 to ABQ72214 and ABP48191 to ABP51230 represent DNA target sequences and zinc finger peptides which are given
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a zinc finger protein (I) that binds to a target site, comprising a first (F1), a second (F2), and a third (F3) zinc finger, ordered F1, F2, F3 from N-terminus to C-terminus, where the target site comprises, in 3'-5' direction, a first (S1), a second (S2), and a third (S3) target subsite, his observibed are: (1) a polypeptide (II) comprising (I); (2) a polynucleotide (III) encoding (I) or (II); and (S3) designing (M) (I) involves selecting the F1 zinc finger such that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New zinc finger protein that binds to target site, useful in studying gene function and for human therapeutics and plant engineering, comprises first, second and third zinc fingers, ordered from N- to
                                                   28-AUG-2002
                                                                                                                                                                                                                                                                                                                                            Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 56; 81pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C-terminus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Liu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-NOV-2000; 2000US-0716637
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zinc finger protein; ZFP; DNA binding protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zinc finger protein related oligonucleotide target SEQ ID NO:2148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-500284/53
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                                                                                                                                                                                                                                                                                                                                                                              exemplification
                                                                                                                                                                                                                                         5 GGGCCAGGC 13
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                                                                                                                                                                                                                                                                          Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SANGAMO BIOSCIENCES INC
                                                                                                                                                                                                           GGGGCAGGC 9
                                                                                                                                                                                                                                                                                                                                            BP; 1
                                                                                                                                                                                                                                                                          Conservative
                                                   (first entry)
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                                                                                                                                                                                                                                                                                         37.0%;
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                                                                                                                                                                                                                                                                                                                                                                            of the present invention
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                                                                                                                                                                                                                                                                                         Score 7.4;
Pred. No. 5
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                                                                                                                                                                                                                                                                                                            DB
                                                                                                                                                                                                                                                                                          2e+02;
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                                                                                                                                                                                                                                                                                                         Length 9;
                                                                                                                                                                                                                                                                          Indels
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                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                             given
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Zinc finger protein related oligonucleotide

target SEQ ID NO:2149

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RESULT 251
AAZ82328
ID AAZ823
XX AAZ822
AC AAZ822
XX 07-APF
XX 07-APF
XX MHUMBAN,
KW HUMBAN,
KW ANTIME
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes a zinc finger protein (I) that binds to C2 a target site, comprising a first (FI), a second (F2), and a third (F3) C2 zinc finger, ordered FI, F2, F3 from N-terminus to C2-terminus, where the carget site comprises, in 3'-5' direction, a first (S1), a second (S2), and a third (S3) target subsite. Also described are: (1) a polypeptide (II) comprising (I); (2) a polymucleotide (III) encoding (I) or (II); and (C2) designing (M) (I) involves selecting the F1 zinc finger such that it binds to the S2 target subsite, selecting the F2 zinc finger such that it binds to the S3 target subsite, and selecting the F3 zinc finger such that it binds to the S3 target subsite, and selecting the F3 zinc (C3) designing (I) is useful for recognition of triplet target subsites having the nucleotide (G1) in this designing (I) that binds to a target site. (I) is useful for recognition of the S3 subsite. (I) is useful in studying gene function, and for human (C4) therapeutics and plant engineering. (I), (II) or (III) is useful in the the expression of a target region within C4 subsite. (I) and diagnostic methods for sequence specific detection of target nucleic acid in a sample, and in assays to determined the phenotype and function of gene expression. (I) has improved affinity and specificity for their target sequences, as well as enhanced conditions of the present DNA target sequences and zinc finger peptides which are given in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local :
Human; metastatic breast tumour tissue; breast cancer; tag; primer; non-metastatic breast tumour tissue; gene therapy; anticancer; antimetastatic; vaccine; diagnosis; ss.
                                                                            Metastatic
                                                                                                                 07-APR-2000
                                                                                                                                                      AAZ82328
                                                                                                                                                                                       AAZ82328 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 9 BP; 1 A; 2 C; 6 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New zinc finger protein that binds to target site, useful in studying gene function and for human therapeutics and plant engineering, comprises first, second and third zinc fingers, ordered from N- to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Liu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 56; 81pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-NOV-2000; 2000US-0716637.
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                                                                            breast
                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                               (first entry)
                                                                      tumour cell upregulated transcript tag #1562
                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                88.98;
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Pred. No. 5.2e+02;
0; Mismatches 1
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RESULT 252
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Best Local S
Matches 8
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19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                          monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for
                             Human dendritic cell SAGE tag, SEQ ID NO:405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells).

AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour
                                                                                           AAZ77977
                                                            10-APR-2000
                                                                                                                        AAZ77977 standard; DNA; 10
                                                                                                                                                                                                                                                                                                                               Sequence 10
                                                                                                                                                                                                                                                                                                                                                              adoptive immunotherapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cells). These transcripts can be used for diagnosis, prognosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 100; 219pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             prevention and treatment of cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-DEC-1999
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(ROBE/)
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8; Conser
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                                                                                                                                                                                                                                    GGCCAGGCC 14
                                                                                                                                                                                                                                                                                                                               BP; 0 A; 4 C; 4
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                                                            (first entry)
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98US-0090039.
98US-0090040.
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88.9%;
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Pred. No. 1.4e+02;
0; Mismatches 1;
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                                                                                                                                                                                                                                                                   Gaps
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SAGE tag; serial analysis of gene expression; antigen-presenting cell;

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with monocytes. Some of the transcripts correspond to known genes or C SSTs (expressed sequence tags) which were previously unknown to be preferentially or differentially expressed in dendritic cells, while cher transcripts correspond to novel genes. Antigen-presenting cell (APC)-associated costimulatory factors play an important role in the activation of the cytotoxic immune response, particularly against tumour cells. Tumour antigen presentation via the MHC (major histocompatibility complex) and subsequent recognition by T-cell receptors is alone in sufficient to activate a robust cytotoxic immune response that can lyse the tumour cells, immunostimulatory cofactors also being required for efficient activation of cytotoxic Tlymphocytes (CTLs). Nucleic acid sequences identified using the SAGE tags have several potential uses. They may be used in vaccines to induce an immune response, particularly against a tumour antigen; to modulate the genotype of an APC; to screen
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19-JUN-1998;
19-JUN-1998;
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19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                    immunostimulatory cofactor proteins which are preferentially or differentially expressed in monocyte-derived dendritic cells compared
                                                                                                                                                                                                                                  Sequences AAZ77573-Z79709 represent SAGE (serial analysis of gene expression) tags used to identify mRNA transcripts encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APC; monocyte-derived dendritic cell; differential gene expression; immunostimulatory cofactor; costimulatory factor; CTL; cytotoxic T-lymphocyte; tumour antigen; immunotherapy; anticancer; ss.
                                                                                                                                                                                                                                                                                                                      Isolated polynucleotides differentially expressed
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19-JUN-1998
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                                                                                                                                                                                                                                                                                                        useful in gene vaccines against cancer
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ROBERTS B L
SHANKARA S.
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98US-0090077.
98US-0090077.
98US-0090078.
98US-0090079.
98US-0090080.
98US-0111715.
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98US-0089993

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98US-0089997

98US-0099000

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AAZ84966
IID AAZ8
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Best Local S
Matches 8
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19-JUN-1998;
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19-JUN-1998;
19-JUN-1998;
AAZ80767 to AAZ83941 represent tags transcripts that are preferentially tumour tissue (i.e. are upregulated AAZ83942 to AAZ86677 represent tags that are preferentially transcribed
                                                                                                                                                                    Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis, prevention and treatment of cancer -
                                                                                                                               Claim
                                                                                                                                                                                                                                                                                                                                                                (GENZ )
(ROBE/)
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ROBERTS E
SHANKARA
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8; Conserv
                                                                                                                            Page 171;
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B L.
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98US-0090040
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                                                                                                                               219pp; English.
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corresponding to distinct transcribed in the metastatic breast in metastatic breast tumour cells). corresponding to distinct transcripts in the primary or non-metastatic
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RESULT 254
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                                                        The invention provides a male tissue-preferred regulatory region (I) comprising nucleotide sequences essential for initiating transcription of the MS45 gene. A method of mediating male fertility in a plant is provided that involves introducing an expression vector comprising a promoter operably linked to (I) into a plant where the exogenous gene impacts male fertility of the plant and (I) controls expression of the exogenous gene. A method of producing hybrid seeds is also provided. Sequences AAH76341-355 represent a series of 5' deletions in the MS45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cents, e.g. cytotoxic T lymphocytes, and these used for
                                                                                                                                                                                                                                                                                                                                                                                                                                         A male tissue-preferred regulatory region comprising nucleotide sequences essential for initiating transcription of the MS45 genuseful for mediating fertility in a male plant -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Albertsen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            monitoring and treatment of breast cancer, particularly where metastandonsis is by standard immunoassays or hybridisation/amplification
                                                                                                                                                                                                                                                                                                                                                                             Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-FEB-2000; 2000US-0504487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-FEB-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zea mays
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hybrid seed;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mays Ms45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAH76342;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 10 BP; 0 A; 3 C; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   breast tumour tissue (i.e. are cells). These transcripts can k
                                   promoter region,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200160997-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     M845;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (PION-) PIONEER HI-BRED INT INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              male tissue; regulatory region; transcription; male
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               N
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8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tumour tissue (i.e. are downregulated in metastatic breast tumour These transcripts can be used for diagnosis, prognosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGCCTGGC 10
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                                                                                                                                                                                                                                                                                                                                                                      Fig 8; 50pp; English.
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88.9%;
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                                   determining
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Huffman
                                      the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        for diagnosis, prognosis, ncer, particularly where metastatic.
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                                                                                                                                                                                                                                                                  The invention provides a male tissue-preferred regulatory region (I) comprising nucleotide sequences essential for initiating transcription of the MS45 gene. A method of mediating male fertility in a plant is provided that involves introducing an expression vector comprising a promoter operably linked to (I) into a plant where the exogenous gene impacts male fertility of the plant and (I) controls expression of the exogenous gene. A method of producing hybrid seeds is also provided. Sequences AAH76341-355 represent a series of 5 deletions in the MS45 promoter region, used for determining the essential region of MS45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A male tissue-preferred regulatory region comprising nucleotide sequences essential for initiating transcription of the MS45 gen
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8; Conser
                                                 CCGGGCCAG 11
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88.9%;
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                                                                                                Score 7.4; DB 1;
Pred. No. 1.4e+02;
0; Mismatches 1
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Pred. No. 1.4e+02;
0; Mismatches 1
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RESULT 257
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Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          comprising nucleotide sequences essential for initiating transcription of the MS45 gene. A method of mediating male fertility in a plant is provided that involves introducing an expression vector comprising a promoter operably linked to (I) into a plant where the exogenous gene impacts male fertility of the plant and (I) controls expression of the exogenous gene. A method of producing hybrid seeds is also provided. Sequences AAH76341-355 represent a series of 5 deletions in the MS45 promoter region, used for determining the essential region of MS45
                                                                                                                                                                                                                                                                                                                        Human; single nucleotide polymorphism; SNP; RANGAP1; haplotyping chromosome 22q13.2-q13.31; Ran GTPase activating protein 1; geno
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A male tissue-preferred regulatory region comprising nucleotide sequences essential for initiating transcription of the MS45 genuseful for mediating fertility in a male plant -
                                                                                                                            WO200179240-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAS19876;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAS19876 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 10 BP; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Albertsen MC,
                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-FEB-2001; 2001WO-US04527
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Z. mays Ms45 promoter deletion mutant fragment LS08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-OCT-2001
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8; Conserv
                                                                                                                                                                                                                                                                                          irregular
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  tissue; regulatory region; transcription; male
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    #56 to detect human RANGAP1 gene polymorphisms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; 5 C; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
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88.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 7.4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                  disorder; primer; ss.
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RESULT 258
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human ubiquitously expressed transcriptome sequence SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Genotyping human Ran GTPase activating protein 1 gene of individual for determining haplotype of individual, involves determining identity of nucleotide pair at specific polymorphic sites for two copies of the gene.
                                                                                                                                                                                                                                                                                                                      24-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                              21-NOV-2000; 2000WO-US31922
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                                                                                                                                                2001-367706/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      transcriptome;
diagnosis; cell
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      specific
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    expression pattern; cancer; drug screening;
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                                                                                                                                                                                                      Kinzler
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Gaps

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New isolated polynucleotides, useful type, such as cancer cell, comprises particular cell types -

comprises

for identifying specific transcriptomes expressed

cell in

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RESULT 259
AAH64375
ID AAH643
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Best Local S
Matches 8
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Best Local Similarity
                                                                                                                                                                                           The present invention describes a method of identifying the type of cell in a sample, involving determining which of the sequences AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce cancer specific gene expression, standardise expression and restore the function of a diseased cell or tissue. The present sequence is one of the transcriptomes described in the exemplification of the invention.
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                                                                                                                                                 Sequence 10 BP; 1 A; 4 C; 3 G; 2 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New isolated polynucleotides, useful for identifying specific type, such as cancer cell, comprises transcriptomes expressed particular cell types -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-367706/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200138577-A2
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8; Conserv
GCCAGGCCA 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 67; 94pp; English.
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                                                  Conservative
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88.9%;
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                                               Score 7.4; DB 1;
Pred. No. 1.4e+02;
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Pred. No. 1.4e+02;
0; Mismatches 1
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RESULT 260
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XX XSOlute
XX Solute
KW Glutam
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Best Local S
Matches 8
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                                Solute carrier family 1; SIC1A4; haplotyping; human; cancer; primer; glutamate/neutral amino acid transporter; neurological disease; PCR;
amino acid transporter disorder; single nucleotide polymorphism; SNP
                                                                                                                            Solute Carrier Family 1 (SLC1A4)
                                                                                                                                                                                          24-SEP-2002
                                                                                                                                                                                                                                                            ABK95840
                                                                                                                                                                                                                                                                                                                      ABK95840 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           encoding AAB98009, which are given in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LPS activated human monocyte expression
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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88.9%;
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Pred. No. 1.4e+02;
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                                                                                                                            primer extension oligonucleotide #11.
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ARBSULT 262
ABV65097/C
ID ABV650
XX ABV650
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XX C ABV650
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XX Human
XX Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC (glutamate/neutral mino acid transporter), member 4 (SICIA4) isogenes CC (II) and an isolated polypeptide (III) comprising an amino acid sequence CC which is a polymorphic variant of a reference sequence for SICIA4 CC protein. Also described are methods for: (I) haplotyping or genotyping SICIA4 gene of an individual; (2) predicting a haplotype pair for SICIA4 CC gene of an individual; (3) identifying an association between a trait and CC at least one haplotype or haplotype pair of SICIA4 gene. (III) Is useful CC in screening for drugs targeting (III) that are useful for treating CC enter, neurological diseases and amino acid transporter disorders. The CC enter thods are useful for improving the efficiency and reliability of CC diseases associated with SICIA4 activity. The haplotyping method is also CC used by the pharmaceutical research scientist to validate SICIA4 as a CC candidate target for treating a specific condition or disease predicted CC trials for treating a specific condition or disease predicted CC and amino acid transporter disorders, and in the design of clinical CC trials for treating a specific condition of disease associated with SICIA4 activity, e.g. cancer, neurological diseases CC and amino acid transporter disorders, and in the design of clinical CC trials for treating a specific condition of disease associated with SICIA4 activity. The methods are also useful in diagnostic, CC prognostic and therapeutic methods. ABK95761-ABK95877 represent CC SICIA4 gene allele-specific coligonucleotides, primer extension CC oligonucleotides and related PCR primers used to identify single CC nucleotide polymorphisms (SNP) of the gene.
                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 8
Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                   Human skin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel genetic variants of Solute Carrier Family 1 (Glutamate/Neutral Amino Acid Transporter), Member 4 isogenes, for improving efficiency and reliability in drug development for treating cancers -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-519580/55
                                                                                                                                                                  ABV65097;
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                                                                                   EST 2883.
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                                                                                                                        (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence which comprises
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                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. 1.4e+02;
); Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
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RESULT 263
ABV69411
ID ABV6944
XX ABV694
XX ABV694
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XX Human
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XX DSOT14
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Best Local S
Matches
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                                                                                                                                                                                           Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                         Human skin EST 7197.
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                                           03-JAN-2001; 2001DE-1000127
                                                                         20-DEC-2001; 2001WO-EP15179
                                                                                                                                    WO200253774-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag
                         The present invention a designed to cleave RNA other than cell-cycle of
                                                                                                                                                                                          New hairpin and hammerhead ribozyme RNA encoding a cyclin or cell-cycle
                                                                                                                                                                                                                                                                                                                                                                                                                     04-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mammalia
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                                                                                                                        Disclosure;
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o cleave RNA encoding a cyclin or c
cell-cycle dependent kinases CDK1,
tive examples of ribozyme recogniti
                      invention relates to a hairpin or hammerhead ribozyme cleave RNA encoding a cyclin or cell-cycle dependent lcell-cycle dependent kinases CDK1, PCNA and Cyclin B1
                                                                                                                      Page
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lin B1 -
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ying cosmetic or pharmaceutical agents against
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                                                                                                                        English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 7.4; DB 1;
Pred. No. 1.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     w
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                                                                                                                                                                                                                                                                                                                   Robbins
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                                                                                                                                                                                            for inhibiting restenosis, cleaves dependent kinase other than CDK1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                   skin or eye disease and scarring. The method involves administering ribozyme (I) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproterinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering nucleic acid molecule (II) comprising a promoter operably linked in nucleic acid segment encoding (I) (I) can have antipsoriatic, dermatological, cytostatic, antiseborrheic, antidiabetic, antisic ophthalmological, vulnerary, keratolytic and virucide activities.
cleaves RNA encoding cytokine involved in inflammation. (I) can be useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keraros.
                                                                                                                                                                                                                    Treating proliferative skin or eye diseases and scarring, ribozymes that cleave RNA encoding cytokines involved in matrix metalloproteinases, growth factors and cell-cycle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytosta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA82415 to AAA86787. The ribozyme of the invention is useful inhibiting restenosis by introduction of the ribozyme into cell The ribozyme is resistant to endonuclease activity and hence i efficient in restenosis treatment.
                                                                                                                                     The present invention skin or eye disease ar
                                                                                                                                                                             Example 1;
                                                                                                                                                                                                                                                                                                        Robbins JM,
                                                                                                                                                                                                                                                                                                                                                             26-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisickling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           recognition site; target; ribozyme binding site; eye disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cell-cycle dependent kinase cdk4 ribozyme binding site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ribozyme therapy; hairpin ribozyme; hammerhead ribozyme;
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                                                                                                                                                                             Page 105; 408pp;
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                                                                                                                                     describes a method for treating nd scarring. The method involves
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Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing scarring such as keloid, adhesion and hypertrophic or hypertrophic burn scarriffication to AAHS2099 represent sequences used in the
                                                                                                      Sequence 8
                                                                                                                                                                                                                     The 5'-primers AAT09358-508, and the 3'-primers AAT09509-659, which target human protein coding regions, together comprise a PCR primer kit with 1361 possible primer pairs. The kit is used in a new method for the characterisation of nucleic acid sequences obtd. from human biological samples, which comprises PCR amplification and indexing of the prods. w.r.t the primer pair that hybridised to its delineating subsequences. The method may be used in the identification, cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Characterisation of nucleotide sequences using primer pairs - by amplification and indexing of amplification prods. w.r.t. primers used for genome mapping and disease diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      characterisation; identification; c.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3'-primer used for characterisation of human biological samples
                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 19; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1996-010958/01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disease diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3'-primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-MAR-2003
25-JUN-1996
                                                                                                                                                                                                and analysis of genes, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-NOV-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lopeznieto
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  Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
8; Conserv
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  Conservative
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(first entry)
                                                                                                    0 A;
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biological samples; PCF
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                                                                                                    3 C; 2 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             37.0%;
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                                                                                                                                                     correct
     0
                                                                                                                                                                                                in genome mapping,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                Score 7;
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Pred. No. 2.4e+02;
                              Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      invention.
     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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                            .DB 1; Lei
o. 5.8e+02;
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PCR amplification; indexing
                                                                                                                                                     field.)
                                                Length 8;
     0,
                                                                                                                                                                                                     and disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
     Indels
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Gaps
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RESULT 267
AATO9656/c
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RESULT 268
AAT09366
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Best Local Similarity

Matches 7; Conserv
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                                                    AAT09366
                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 8
                                                                                                   AAT09366 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and analysis of genes,
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25-JUN-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    identification; cloning; analysis; genes; genome mapping;
disease diagnosis; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                 B₽;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (updated)
(first entry)
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                                                                                                     ВP
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                                                                                                                                                                                                                                                                                                                                                                                  Score 7;
                                                                                                                                                                                                                                                                                                                                   Pred. No. 5. 
Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                    other;
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o. 5.8e+02;
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25-MAR-2003

(updated)

21-JUN-1996

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RESULT 269
AAT09389
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Best Local S
Matches 7
                                                                     5'-primer; human; protein coding region; PCR primer kit;
characterisation; biological samples; PCR amplification; indexing
identification; cloning; analysis; genes; genome mapping;
                                                                                                                       5'-primer used for characterisation of human
                                                                                                                                                                                                                                                                                                                                                                                                                                     The 5'-primers AAT09358-508, and the 3'-primers AAT09509-659, which target human protein coding regions, together comprise a PCR primer kit with 1361 possible primer pairs. The kit is used in a new method for the characterisation of nucleic acid sequences obtd. from human biological samples, which comprises PCR amplification and indexing of the prods. w.r.t the primer pair that hybridised to its delineating subsequences. The method may be used in the identification, cloning
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Characterisation of nucleotide sequences using primer pairs - by amplification and indexing of amplification prods. w.r.t. primers used for genome mapping and disease diagnosis
           WO9531574-A1
                                    Synthetic
                                                          disease
                                                                                                                                                 25-MAR-2003
21-JUN-1996
                                                                                                                                                                                                              AAT09389 standard; DNA; 8
                                                                                                                                                                                                                                                                                                                                                                             Sequence 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5'-primer used
                                                                                                                                                                                       AAT09389;
                                                                                                                                                                                                                                                                                                                                                                                                                          and analysis of genes, e.g. in genome mapping, and disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1996-010958/01
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   characterisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lopeznieto
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                                                                                                                                                                                                                                                                            N
                                                          diagnosis;
                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                   AGGCCAA 16
                                                                                                                                                                                                                                                                                                                                                                                                     on 25-MAR-2003 to
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human; protein coding region; PCR primer kit;
sation; biological samples; PCR amplification; indexing;
                                                                                                                                                                                                                                                                                                                                                                             BP; 4 A;
                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                               (updated)
(first entry)
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                                                                                                                        biological samples
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                                                                                                                                                                                                                                   5'-primer; human; protein coding region; PCR primer kit; characterisation; biological samples; PCR amplification; identification; cloning; analysis; genes; genome mapping;
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                         WPI; 1996-010958/01
                                                                                                                         12-MAY-1995;
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                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                        disease diagnosis; ss
                                                                                                                                                                                                                                                                                      5'-primer used for characterisation of human biological samples
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21-JUN-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Characterisation of nucleotide sequences using primer pairs - by PCR amplification and indexing of amplification prods. w.r.t. primers used for genome mapping and disease diagnosis
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Characterisation of nucleotide sequences using primer pairs -

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RESULT 271
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XX Synth
XX 23-NO'
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                  The 5'-primers AAT09358-508, and the 3'-primers AAT09509-659, which target human protein coding regions, together comprise a PCR primer kit with 1361 possible primer pairs. The kit is used in a new method for the characterisation of nucleic acid sequences obtd. from human biological samples, which comprises PCR amplification and indexing of the prods. w.r.t the primer pair that hybridised to its delineating subsequences. The method may be used in the identification, cloning
                                                                                                                                                                                                                             Disclosure; Page 19;
                                                                                                                                                                                                                                                                                                 Characterisation of nucleotide sequences using primer pairs - by amplification and indexing of amplification prods. w.r.t. primers
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25-JUN-1996
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of genes,
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                                                                                                                                                                                                                                                                         mapping and
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biological samples; PCR amplification;
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XX AAX295
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DE Primer
XX Nuclei
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OS Synthe
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                                                                                                                                                                                in plant biology, agriculture, human or veterinary medicine, reproduction, microbiology or environmental science, e.g. to study expression of nuclear receptors at different stages of tissue deve or after treatment with particular drugs. It is also used for DNA fingerprinting (to generate products useful for differential hybridisation), or, where a 3'-anchor primer is used, to isolate the stage of mRNA sequences. Sequences AAX29526-X29583 represent cla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amplification; nuclear receptor; G-protein coupled receptor; apoptosis; DNA repair; DNA replication; plant biology; agriculture; human; veterinary medicine; reproduction; microbiology; hybridisation; environmental science; DNA fingerprinting: PCP primar. --
                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                   The invention provides primers (AAX29501-X29679) for identifying sequences encoding structurally or functionally related proteins such as nuclear or G-protein coupled receptors, apoptosis-related or DNA repair/replication proteins. The identified sequences are broadly useful
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(Updated on 25-MAR-2003 to
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7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                       primers able to amplify
                                                                                                                                   BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP;
                                                                  Conservative
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                                                                                                                                   other;
                                                                                                                                                                  coupled receptor genes
                                                                                   DB 1; Le
b. 5.8e+02;
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5.8e+02;
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RESULT 274
AAA80793
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AAX29558/c
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                                                                                                                                                                                                                                                                                                         맑
                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                                                                                                                                                                              reproduction, microbiology or environmental science, e.g. to study expression of nuclear receptors at different stages of tissue development or after treatment with particular drugs. It is also used for DNA fingerprinting (to generate products useful for differential hybridisation), or, where a 3'-anchor primer is used, to isolate the 3'-ends of mRNA sequences. Sequences AAX29526-X29583 represent claimed primers specific for human G-protein coupled receptor genes.
                                    A. thaliana primer walking octamer SEQ ID NO: 106.
                                                                                                                                                                                                                                                                                                       Sequence 8
                                                                                                                                                                                                                                                                                                                                                                                                                                nuclear or G-protein coupled receptors, apoptosis-related or DNA repair/replication proteins. The identified sequences are broadly useful in plant biology, agriculture, human or veterinary medicine,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention provides primers (AAX29501-X29679) for identifying sequences encoding structurally or functionally related proteins
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Subset of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nucleic acid amplification; nuclear receptor; G-protein coupled receptor; apoptosis; DNA repair; DNA replication; plant biology; agriculture; human; veterinary medicine; reproduction; microbiology; hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAX29558 standard; DNA; 8
          Primer walking; octamer; primer; DNA sequencing; PCR;
                                                                 24-NOV-2000
                                                                                             AAA80793;
                                                                                                                        AAA80793 standard; DNA; 8 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-205200/17.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         04-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  environmental science; DNA fingerprinting; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Primer for human G-protein coupled receptor genes.
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                                                                                                                                                                                           GGCCAGG 1
                                                                                                                                                                                                                      GGCCAGG 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                primers able to amplify group of related sequences
                                                                                                                                                                                                                                                                                                       BP; 0 A;
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100.0%; Pr/
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Best Local S
Matches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       molecule which comprises selecting a library primer from an octamer oligonucleotide library consisting of 48 bp sequences and corresponding complementary sequences, where the library primer is complementary to a known sequence adjacent to the unknown sequence or is complementary to a sequence in a known extension product. The method is useful for DNA nucleotide sequencing, in PCR, and in other processes which make use of primers. The octamers are used to identify coding sequences. Primer walking using the octamer libraries is advantageous over other sequencing methods because it does not require multiple cloning steps nor subsequent template preparations, and it is a directed and methodical approach. AAA80688 A81253 represent the octamer primers used in the primer walking method of the invention.
                                                                                                                                 US6083695-A
                                                                                                                                                                                                             A. thaliana primer walking octamer SEQ ID NO: 359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 8 BP; 4 A; 2 C; 2 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and other primer processes oligonucleotides -
                                                                             21-MAY-1997;
                                                                                                       04-JUL-2000
                                                                                                                                                          Arabidopsis thaliana
                                                                                                                                                                                 Primer walking; octamer; primer; DNA sequencing; PCR;
                                                                                                                                                                                                                                         24-NOV-2000
                                                                                                                                                                                                                                                                 AAA81046;
                                                                                                                                                                                                                                                                                            AAA81046 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequencing an unknown DNA molecule for the polymerase chain reaction and other primer processes comprises primer walking of octamer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hardin PE, Hardin SH,
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             (UYHO-) UNIV HOUSTON (HARD/) HARDIN S H.
                                                   15-APR-1996;
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                                                                             97US-0859954
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5.8e+02;
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RESULT 276
ADD22387/c
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes a novel method for sequencing an unknown DNA molecule which comprises selecting a library primer from an octamer oligonucleotide library consisting of 48 8-bp sequences and corresponding complementary sequences, where the library primer is complementary to a known sequence adjacent to the unknown sequence or complementary to a known sequence in a known extension product. The method
                                                                                                                          Modifying endogenous oil of cotton plants, to produce cotton seed with reduced palmitic and/or linoleic acid content, involves product transgenic plants containing a fatty acid biosynthesis gene in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequencing an unknown DNA molecule for the polymerase chain and other primer processes comprises primer walking of octamoligonucleotides -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cotton; delta9-desaturase; fatty acid delta12-desaturase; palmitic linoleic acid; stearic acid; oleic acid; transgenic plant;
                                                   Example 11; Page
                                                                                                          construct
                                                                                                                                                                                                                                        WPI; 2001-602932/68
                                                                                                                                                                                                                                                                                                                                                                                                 18-APR-2000; 2000US-198124P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               seed oil; ss.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                             ORG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; L. No. 5.8e+02; 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              octamer
                                                                                                                                s producing in a
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invention

relates

to a method for modifying the endogenous oil

o f

expression and

or

activity of target polypeptides associated

down-regulate the ociated with

The present invention describes low adenosine (A) content antisense oligonucleotides and compositions (I) comprising them. In the antisense oligonucleotides the A is replaced by a 'Universal' or alternative base. (I) can have respiratory, bronchodilator, antiinflammatory, analgesic, immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. The antisense oligonucleotides and (I) can be used to down-regulate the

Claim 14;

Page 106; 1592pp; English respiratory obstructions

and

Low adenosine (A) content antisense oligonucleotides which trigger adenosine receptors during metabolism, useful e.g.

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for treating

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RESULT 277
AAF20849/C
ID AAF208
XX AAF208
AC AAF208
XX Low ad
XW Low ad
XW human;
XW respir
XW respir
XW surfac
XW respir
XW cancer
XW Cancer
XW Cancer
XX Cancer
XX Cancer
XX CACC
XX OC-APR
XX (UYEC-PA (NYCE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             밁
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Best Local S
Matches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Low adenosine antisense oligonucleotide; phosphorothioate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic; respiratory; obstruction; pulmonary obstruction; impeded respiration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             desaturase) gene, or fatty acid delta12-desaturase (oleoyl-PC delta12-desaturase) gene in the seed of a cotton plant. The invention is useful for producing cottonseed oil with reduced palmitic and/or linoleic acid content, and increased stearic and/or oleic acid content. The present DNA sequence is an oligonucleotide which is used for generating fatty acid biosynthesis gene construct related to the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (NYCE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pain; cystic fibrosis; allergic rhinitis; pulmonary hypertension; emphysema; pulmonary transplantation rejection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    a transgenic cotton plant having a gene construct which includes a fat acid biosynthesis gene operably linked to a promoter sequence capable conferring expression of the delta9-desaturase (delta9 stearoyl-ACP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-MAR-2000; 2000WO-US08020
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human adenosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF20849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF20849 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 8 BP; 0 A; 3 C; 5 G; 0 U; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cotton plant, to produce cotton seed oil. The method comprises producing
                                                                                                                                                                                                                                                                                                                                                                                                          2000-679539/66.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            UNIV EAST CAROLINA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                obstructive pulmonary disease; pulmonary infection; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ccggccc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-0127958
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Al receptor polynucleotide fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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RESULT 278
AAA34727/C
ID AAA347
XX AAA347
XX AAA347
XX AAA347
XX Human
XX Cancer
XX Cancer
XX Pulmon
XX WO2000
XX WO2000
XX Cancer
XX UYEC-
XX UYEC-
XX WPI;

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Best Local S
Matches 7
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                             New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstriction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                phosphorothioate; impaired respiration; inflammation; allergy; allergic disease; bronchoconstriction; inhibitor; antiinflammatory; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; lung disease; ischaemic condition; pulmonary vasoconstriction; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            condition selected from pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome (RDS), pain, cystic fibrosis (CF), allergic rhinitis (AR), pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                lung/respiratory disorders and malignancies, such as stimulating and activating peptide factors and transmitters, transcription factors, immunoglobulins and antibodies, antibody receptors, cytokines and chemokines, endogenously produced specific and non-specific enzymes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pulmonary transplantation rejection, pulmonary infections, bronchitis, and/or cancer. AAF18434 to AAF21543 represent human polynucleotide fragments and antisense oligonucleotides used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200009525-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       respiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; COPD.
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                                                                                                                                                                                                        WPI; 2000-205971/18.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; adenosine receptor; low adenosine antisense oligonucleotide;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA34727;
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                                                                                                                                                                                                                                                                                                                                                                           (UYEC-) UNIV EAST CAROLINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 leukaemia;
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3
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100.0%; Pr
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k; Pred. No. 5.2
0; Mismatches
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o. 5.2e+02;
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Query Match
Best Local Similarity
7; Conserve
                                                                                                                                                                         fibrosis, pulmonary hypertension, emphysema, chronic obstructive (CC pulmonary disease (COPD), and cancers such as leukaemias, lymphomas, CC carcinomas, and cancers which may metastasise to the lungs, including the breast and prostate cancer. The reduction of the adenosine content of the ONs reduces side effects. The A-containing ONs break down with the CC release of deoxyadenosine which activates adenosine receptors causing tronchoconstriction and inflammation. AAA32313 to AAA35312 represent the nucleotide sequences given in the sequence listing from the present the invention, which correspond to SEQ ID NO:1 to 2815, and then the last CC invention, which correspond to SEQ ID NO:1 to 185, but the sequences CC (AAA32312 to AAA33912) are specifically claimed ONs from the present correspond to AAA33912 represent the invention. N. B. Sequences given in the disclosure of the present corresponding SEQ ID NO: sequences given in the disclosure of the present convention do not match up with their corresponding SEQ ID NO: sequences yver in the disclosure of the present corresponding SEQ ID NO: sequences yver in the disclosure of the present corresponding SEQ ID NO: sequences yver in the disclosure of the present corresponding SEQ ID NO: sequences yver in the disclosure of the present corresponding SEQ ID NO: sequences yver in the sequences sequences of the present corresponding SEQ ID NO: sequences yver in the disclosure of the present corresponding SEQ ID NO: sequences yver in the sequences sequences of the present corresponding SEQ ID NO: sequences yver in the disclosure of the present corresponding SEQ ID NO: sequences yver in the sequences of the present corresponding SEQ ID NO: sequences yver in the sequences of the present corresponding SEQ ID NO: sequences yver in the sequences of the present corresponding SEQ ID NO: sequences yver in the sequences of the present corresponding SEQ ID NO: sequences yver in the sequences of the present corresponding SEQ ID NO: sequences of the present corresponding SEQ ID NO: sequen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antisense oligonucleotide (ON) with low adenosine (up to 15%), which targets nucleic acids involved in bronchoconstriction, allergies, and/or inflammation. The ON can have antiinflammatory, antialtergic, antiasthmatic, cytostatic and analgesic activities. The compositions are useful for the treatment of diseases associated with inflammation, eighbarred altways, including lung diseases and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating
                                                                                                                                                                     Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ischaemic conditions, pulmonary vasoconstriction, allergies, na, impeded respiration, respiratory distress syndrome, pain,
                                                                                                                                                                  BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 571; 1343pp; English
                                    Conservative
                                                                                                                                                                  A; 3 C; 5 G; 1 T;
                  35.0%; 5--
100.0%; Prr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 describes a new
                                Score 7; DB 1; Pred. No. 5.2 0; Mismatches
                                                                                                                                                                         0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         composition comprising an ow adenosine (up to 15%), whincheconstriction, allergies,
                                                                DB 1; Le:
5.2e+02;
                                                                                             Length 9;
                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      pain, cystic
                                    0
                                    Gaps
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                                                                                                                                                                                                                                                                                                                                 RESULT 279
        New zinc finger protein that binds to target site, useful in studying gene function and for human therapeutics and plant engineering, comprises first, second and third zinc fingers, ordered from N- to
                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                               Zinc finger
                                                                                                                                                                                                                                                                    28-AUG-2002
                                                                                                                                                                                                                                                                                        ABQ72098
                                                                                                                                                                                                                                                                                                             ABQ72098 standard; DNA; 9 BP
                                                 WPI; 2002-500284/53.
                                                                                                             20-NOV-2000; 2000US-0716637
                                                                                                                                  20-NOV-2001; 2001WO-US43438
                                                                                                                                                      30-MAY-2002
                                                                                                                                                                          WO200242459-A2
                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                          Zinc finger protein; ZFP; DNA binding protein; zinc finger; ss
                                                                                          (SANG-) SANGAMO BIOSCIENCES INC
                                                                                                                                                                                                                                                                                                                                                              9
                                                                                                                                                                                                                                                                                                                                                                                  æ
                                                                                                                                                                                                                                                                                                                                                                                  CCAGGCC 14
                                                                                                                                                                                                                                               protein related oligonucleotide target SEQ ID NO:2396
                                                                                                                                                                                                                                                                   (first
                                                                                                                                                                                                                                                                   entry)
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BE CONTROL OF THE CON
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CC a target site, comprising a first (F1), a second (F2), and a third (F3) (C zinc finger, ordered F1, F2, F3 from N-terminus to C-terminus, where the CC target site comprises, in 3'-5' direction, a first (S1), a second (S2), and a third (S3) target subsite. Also described are: (1) a polypeptide (C (II) comprising (I); (2) a polyprucleotide (III) encoding (I) or (II); and (C (3) designing (M) (I) involves selecting the F1 zinc finger such that CC that it binds to the S1 target subsite, selecting the F2 zinc finger such that (C (1) that it binds to the S2 target subsite, and selecting the F3 zinc (I) that binds to a target site. (I) is useful for recognition of triplet (C (1) that binds to a target site. (I) is useful for recognition of triplet (I) that binds to a target site. (I) is useful for recognition of the C (1) is useful in studying gene function, and for human (C (1) therapeutics mathods to modulate the expression of a target region within (C (1) the subsite and plant engineering. (I). (II) or (III) is useful in the S2 target nucleic acid in a smaple, and in assays to determined the (C (2) phenotype and function of gene expression. (I) has improved affinity (C (2) phenotype and function of gene expression. (I) has improved affinity (1) singulated the subsideal activity. ABO71213 to ABO72214 and ABP48191 to ABB51230 (1) the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                     New zinc finger protein that binds to target site, useful in studying gene function and for human therapeutics and plant engineering, comprises first, second and third zinc fingers, ordered from N- to
                                                                                                                                                                                                                                                    WPI; 2002-500284/53.
                                                                                                                                                                                                                                                                                                                  ntJ
                                                                                                                                                                                                                                                                                                                                                                                                                                       20-NOV-2000; 2000US-0716637
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-NOV-2001; 2001WO-US43438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-MAY-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Zinc finger protein; ZFP; DNA binding protein; zinc finger; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zinc finger protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABQ72099;
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                                                                                                                                                                                                                                                                                                                                                                                (SANG-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200242459-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABQ72099 standard; DNA; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 61; 81pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12
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                                                                                                                                                                                                                                                                                                                                                                                SANGAMO BIOSCIENCES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GCCAAAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GCCAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention describes a zinc finger protein (I) that binds to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     related oligonucleotide target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              35.0%;
100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 G; 3 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              %; Score 7; DB 1
%; Pred. No. 5.2
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 No. 5.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO: 2397.
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Example 1;

Page

61;

81pp; English

The invention relates to an isolated polynucleotide (I) comprising a first nucleotide sequence which comprises solute carrier family 1 (glutamate/neutral amino acid transporter), member 4 (SLC1A4) isogenes

Novel genetic variants of Solute Carrier Family 1 (Glutamate/Neutral Amino Acid Transporter), Member 4 isogenes, for improving efficiency and reliability in drug development for treating cancers -

WPI; 2002-519580/55

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Kazemi A,

Russo

DP,

Sausker

30-NOV-2000; 2000US-250254P 29-NOV-2001; 2001WO-US44781

GENAISSANCE PHARM INC.

Claim 17; Page 16; 139pp; English.

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                                                                                                                                                          S
                                                                                                                                                                                                                                                                                                     cc a target site, comprising a first (FI), a second (F2), and a third (F3) crinc finger, ordered F1, F2, F3 from N-terminus to C-terminus, where the ctarget site comprises, in 3'-5' direction, a first (S1), a second (S2), cc and a third (S3) target subsite. Also described are: (1) a polypeptide (II) comprising (I); (2) a polypurelectide (III) encoding (I) or (II); and (C3) designing (M) (I) involves selecting the F1 zinc finger such that (C3) thinds to the S1 target subsite, selecting the F2 zinc finger such that (C4) that it binds to the S2 target subsite, and selecting the F3 zinc (C5) that it binds to the S2 target subsite, and selecting the F3 zinc (C4) that binds to a target site. (I) is useful for recognition of triplet (C5) that binds to a target site. (I) is useful for recognition of triplet (C5) that binds to a target site. (I) is useful for recognition of the C5 subsite. (I) is useful in studying gene function, and for human (C5) therapeutics and plant engineering. (I), (II) or (III) is useful in the subject, in diagnostic methods for sequence specific detection of target nucleic acid in a sample, and in assays to determined the complex to the subsite of the sequence, as well as enhanced (C5) biological activity. ABO71213 to ABO72214 and ABP84891 to ABP51230 (C7) in the exemplification of the present invention.
                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                              Query Match
Best Local
ABK95837 standard; DNA; 10
                                                                                                                                                                                                                                                                                                     Sequence 9 BP; 1 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes a zinc finger protein (I) that binds to
                                                                                                                                                            12 GCCAAAG 18
                                                                                                                œ
                                                                                                                                                                                                        Similarity 7; Conserv
                                                                                                                GCCAAAG 2
                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                     2 C;
                                                                                                                                                                                                        35.0%; Score 7; DB 1
100.0%; Pred. No. 5.2
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                     w
                                                                                                                                                                                                                                                                                                     G; 3 T; 0 other;
                                                                                                                                                                                                                              DB 1; Ler
vo. 5.2e+02;
                                                                                                                                                                                                                                                       Length 9;
                                                                                                                                                                                                             Indels
                                                                                                                                                                                                             0
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RESULT 281
ABK95837/c
ID ABK958
XX ABK958
XX Solute
XX MO2002
XX MO91; 2
XX Novel
PT Amino
PT Amino
CC First
CC (gluta
                                                                                                                                                                                                                                                                                                                                                                                                                         Solute carrier family 1; SLC1A4; haplotyping; human; cancer; primer; glutamate/neutral amino acid transporter; neurological disease; PCR; amino acid transporter disorder; single nucleotide polymorphism; SNP
                                                                                                                                                                                                                                                                                                                            06-JUN-2002
                                                                                                                                                                                                                                                                                                                                                           WO200244198-A2
                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Solute Carrier Family 1 (SLC1A4) primer extension oligonucleotide #8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABK95837;
                                                                                                                                                                                                                                                                                                                                                                                                                      acid transporter disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВР
                                                                                                                                                                                                                                                                                                                                                                                                                        polymorphism;
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RESULT 282
ABV63037/c
ID ABV630
XX ABV630
XX Human
AC Human
XX Human
XX Human
XX Homo s
PN W02002
XX Homo s
PN W02002
XX Homo s
XX Homo s
YX IN UNCLOSE
YX IN Vit
PT Homeos
YX Discl
XX Discl
XX Discl
XX Discl
XX The in
CC in th
CC in th
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   which is a polymorphic variant of a reference semierra for sicilar and a serious and serio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                 In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-JAN-2001; 2001DE-1000127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (HENK ) HENKEL KGAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABV63037 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                    2002-590638/63
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                                                                                                                                                                                                                                         cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EST 823
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
A;
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100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hofmann
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Pred. No.
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The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE)

Query Match

Sequence 11

BP; 4 A; 4 C; 2 G; 1 T; 0 other;

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Length 11;

Disclosure; Page 48;

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RESULT 283
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                                                                                   The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; sclerodermatichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against e.g. skin cancer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-DEC-2001; 2001WO-EP15179
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                                                             The present sequence is that of the invention.
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RESULT 284
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RESULT 285
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Matches
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Matches 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     useful for: identifying markers of skin ageing and/or stress; determining skin ageing and/or stress; and identifying or determining the effects of pharmaceutical or cosmetic agents for control of skin ageing. The present sequence is one of a group of human skin ageing/stress related expressed sequence tags (ABQ86246-ABQ87680) of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            or animals, are important for skin ageing and/or skin stress by serial analysis of gene expression between mixtures of transcribed and optionally translated, genetically encoded factors (A) obtained from young and aged skin, to identify that genes that show strong differential expression. (A) comprises protein or mRNAs or their fragments. (M1) is
                           08-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Identifying genes involved screening for cosmetic or t gene expression -
                                                                            AAS19877;
                                                                                                                       AAS19877 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 11 BP; 4 A; 4 C; 2 G; 1 T; 0
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                                                                                                                                                                                                                                                                                                                                                                          100.0%;
                                                                                                                                                                                                                                                                                                                                                                                              35.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             lved in skin stress and or therapeutic agents,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   stress;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hofmann
                                                                                                                         ВР
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                                                                                                                                                                                                                                                                                                                                                                     Score 7; I
Pred. No.
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                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                               other;
                                                                                                                                                                                                                                                                                                                                                                        DB 1;
b. 2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                expressed sequence tag; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ageing, useful e.g. in based on differential
                                                                                                                                                                                                                                                                                                                                              0,:
                                                                                                                                                                                                                                                                                                                                                                                         Length 11;
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                                                                                                                                                                                                                                                                                                                                              Gaps
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Query Match
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                                                                                                                            The present invention relates to novel single nucleotide polymorphisms (SNPs) in the human Ran GTPase activating protein I (RANGAPI) gene located on chromosome 229[3] 2-[3] 31, and methods for haplotyping and/or genotyping the RANGAPI gene. The methods of the invention make use of allele-specific oligonucleotides (ASOS) as probes and primers and/or primer-extension oligonucleotides for detecting the RANGAPI gene polymorphisms. The polymucleotides for detecting the RANGAPI gene treatment of diseases associated with RANGAPI activity, such as cancer and other disorders associated with an irregular cell cycle. AAS19891-AAS19898 represent primer-extension oligonucleotides for detecting human RANGAPI gene polymorphisms.
                                                                                                                                                                                                                                                                                                                                                                                             Genotyping human Ran GTPase activating protein 1 gene of individual determining haplotype of individual, involves determining identity of the mucleotide pair at specific polymorphic sites for two copies of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; single nucleotide polymorphism; SNP; RANGAP1; haplotyping chromosome 22q13.2-q13.31; Ran GTPase activating protein 1; genocancer; irregular cell cycle associated disorder; primer; ss.
                                                                                                                          Sequence 10 BP; 0 A; 5 C; 3
                                                                                                                                                                                                                                                                                                                                                    Claim 17; Page 16; 148pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-075068/10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide #57 to detect human RANGAP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-APR-2000; 2000US-198072P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-APR-2001; 2001WO-US12455.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GENA-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENAISSANCE PHARM INC
                               TCCGGGCCAG
                                                             Conservative
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                                                                            34.0%;
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                                                                                                                          G; 2 T; 0 other;
                                                                            Score 6.8; DB 1;
Pred. No. 1.8e+02;
                                                               Mismatches
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                                                                                         Length 10
                                                               Indels
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                                                           Gaps
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RESULT 286
AAX18625
Synthetic. Rattus sp.
                                                               p53; serial analysis of gene expression; SAGE tag; cancer; rat embryo fibroblast; REF; tumour suppressor; cell cycle tumourigenesis; diagnosis; ss.
                                                                                                 p53 serial
                                                                                                                 06-MAY-1999
                                                                                                                                                 AAX18625 standard; DNA; 10
                      WO9901581-A1
                                                                                                 analysis
                                                                                                                 (first
                                                                                               of gene expression
                                                                                                                entry)
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neoplastic;

14-JAN-1999

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RESULT 287
AAZ79322
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Matches 8
 19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
19-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                     AAZ79322 standard;
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                                                                                                                                             W09965924-A2
                                                                                                                                                                                                          cytotoxic T-lymphocyte; tumour antigen;
                                                                                                                                                                                                                                                                                        Human dendritic
                                                                                                                                                                                                                                                                                                                        10-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-106079/09
                                                                                 18-JUN-1999;
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                                                                                                                                                                                                                    SAGE tag; serial analysis of gene expression; antigen-presenting cell; APC; monocyte-derived dendritic cell; differential gene expression; immunostimulatory cofactor; costimulatory factor; CTL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tumourigenesis.
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                                                                                                                                                                                                                                                                                                                    (first entry)
98US-0089833.
98US-0089844.
98US-0089853.
98US-0089878.
                                                                               99WO-US13800
                                                                                                                                                                                                                                                                                   cell SAGE tag,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97US-0051573
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98WO-US13903
                                                                                                                                                                                                                                                                                                                                                                                     DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 6.8;
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                                                                                                                                                                                                                                                                                        SEQ ID NO:1750
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 No.
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                                                                                                                                                                                                        immunotherapy; anticancer; ss
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n two tissue
f gene
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CC (APC) - associated costimulatory factors play an important role in the CC activation of the cytotoxic immune response, particularly against the cells. Tumour antigen presentation via the MHC (major histocompatibility complex) and subsequent recognition by T-cell receptors is alone complex) and subsequent recognition by T-cell receptors is alone complex) and subsequent recognition by T-cell receptors is alone complex) and the AMC (major histocompatibility complex) are response that can complex in the subsequent recognition by T-cell receptors is alone complex) and the subsequent recognition by T-cell receptors is alone complex) and the trivation of cytotoxic T-lymphocytes (CTLs). Nucleic acid sequences identified using the SAGE tags have several potential uses. They may be used in vaccines to induce an immune response, particularly cagainst a tumour antigen; to modulate the genotype of an APC; to screen complex that modulate expression of differentially expressed genes in the APC; and as hybridisation probes/amplification primers for the coding receptors and substitution of the dendritic cell cappession of these genes. Detection of the dendritic cells containing the separation of their encoded proteins, can be used to identify cells as belonging to the monocyte lineage. Cells containing them are used in gene therapy. (or to stimulate components and APC-associated costimulatory factors ensures adequate antigen presentation of co-stimulatory signals, migration to T cells; containing the containing the containing the costimulatory factors ensures adequate antigen presentation of co-stimulatory signals, migration of chemokines for the correspondent of the mare used in gene therapy. (or co-administration of the costimulatory signals, migration of chemokines for the correspondent of the approach factors and secretion of chemokines for the correspondent of the cost of the co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        with monocytes. Some of the transcripts correspond to known genes or ESTs (expressed sequence tags) which were previously unknown to be preferentially or differentially expressed in dendritic cells, while other transcripts correspond to novel genes. Antigen-presenting cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          differentially expressed in monocyte-derived dendritic cells compared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  expression)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GENZ )
(ROBE/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           expression) tags used to identify mRNA transcripts encoding immunostimulatory cofactor proteins which are preferentially
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ROBERTS B
SHANKARA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BL,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAZ77573-Z79709 represent SAGE (serial analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Shankara
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98US-0090048.
98US-0090072.
98US-0090076.
98US-0090077.
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98US-0090036.
98US-0090040.
98US-0090040.
98US-0090041.
98US-0090042.
98US-0090043.
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98US-0089997.
98US-0089999.
98US-0090000.
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98US-0089992.
98US-0089993.
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Sequence 10

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RESULT 288
AAZ83064/c
ID AAZ830
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Best Local
      reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in select cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for ratsing specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be use to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for
                                                                                                                                                                               AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; metastatic breast tumour tissue; breast cancer; tag; p: non-metastatic breast tumour tissue; gene therapy; anticancer
                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 121; 219pp; English.
                                                                                                                                                                                                                                                                                                                                                                                         Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9965928-A2
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                                                                                                                                                                                                                                                                                                                                                                            prevention
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ROBERTS B L
SHANKARA S.
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8; Conserv
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98US-0089997.
98US-0090039.
98US-0090040.
98US-0090041.
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Pred. No. 1.8e+02;
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                                             AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells). CC that are preferentially transcribed in metastatic breast tumour cells). CC AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts CC that are preferentially transcribed in the primary or non-metastatic breast tumour cissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, compitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification creations. Compounds that modulate expression of the transcripts are cotentially useful for treatment of (metastatic) breast cancer, while cotentially useful for treatment of (metastatic) breast cancer, while cotentially expected constructions of the transcripts are used to direct expression, in selected corporates, of e.g. therapeutic genes (also ribozymes or antisense cotentially anativally an antisen-encoding sequence for use in gene or cell-based vaccines, for diagnosis breast cancer and for raising the product of the transcripts are also useful in vaccines; for diagnosis breast cancer and for raising the constitution of the transcripts are also as for a producer transcripts are also as for a producer transcripts are also as for a producer transcripts.
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Best Local S
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19-JUN-1998;
19-JUN-1998;
 specific antibodies (Ab). Ab are used to detect the polypeptides therapeutic agents. Host cells that produce the polypeptides can to expand and isolate populations of educated, antigen-specific i
                                                                                                                                                                                                                                                                                                                                                                                 Isolated polynucleotides differentially expressed between metastatic and non-metastatic breast cancer cells, useful for diagnosis, prevention and treatment of cancer -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          non-metastatic breast tumour tissue; gene therapy; anticancer; antimetastatic; vaccine; diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                      Claim
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ROBERTS E
SHANKARA
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Pred. No. 1.8e+02;
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therapeutic agents.

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RESULT 290
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Matches
                                                                                                        AAZ80767 to AAZ83941 represent tags corresponding to distinct transcripts that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour cells).

AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, progness, monitoring and treatment of breast cancer, particularly where metastatics.
        piagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in select cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising
                                                                                                                                                                                                                                                       Isolated polynucleotides differentially and non-metastatic breast cancer cells, prevention and treatment of cancer -
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19-JUN-1998;
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(SHAN/)
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ROBERTS
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                                                                                                                                                                                                                               Page 181;
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B L.
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14
(Ab). Ab are used to
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ne; diagnosis; ss.
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1.8e+02;
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RESULT 291
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                                                                                                                                                                                                                                                                                                           The present invention describes a method of identifying the type of cell in a sample, involving determining which of the sequences AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce
                                                                                                                                                                                           Sequence 10
                                                                                                                                                                                                                                    cancer specific gene expression, standardise expression and restore t function of a diseased cell or tissue. The present sequence is one of the transcriptomes described in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              type, such as cancer cell, comprises particular cell types -
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 13; Page 59; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated polynucleotides, useful
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UYJO ) UNIV JOHNS HOPKINS
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diagnosis; cell
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l specific gene expression; ss.
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                                                                                                       Score 6.8; DB 1,
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Pred. No. 1.
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                                                                                                                                           Length 10
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                                                                                                                                                                                                                                                                                           restore the
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RESULT 292 AAH64037

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Best Local S
Matches 8
                     respiratory distress syndrome; pain; cystic fibrosis; pulmonary hypertension; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer; hepatocellular carcinoma; kidney cancer; melanoma; hepatic metastäsis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes a method of identifying the type of cell in a sample, involving determining which of the sequences
AAH63161-AAH64724 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce
                                                                                                                                                                                                      Antisense oligonucleotide; multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vasoconstriction; inflammation; allergic rhinitis;
                                                                                                                                                                                                                                                                                                                           C/EBP-beta antisense oligonucleotide fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAX55034 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 10 BP; 1 A; 4 C; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cancer specific gene expression, standardise expression and restore the function of a diseased cell or tissue. The present sequence is one of the transcriptomes described in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated polynucleotides, useful type, such as cancer cell, comprises particular cell types -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Velculescu VE,
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                                                                                                                                                                             acute asthma; allergy; asthma; impeded respiration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-NOV-2000; 2000WO-US31922
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Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The specification describes antisense oligonucleotides (AAX52869-X55271) directed against at least 2 mRNAs selected from target genes, coding and non-coding regions of RNAs corresponding to target genes, gene initiation codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'-end and the juxta-section between coding and non-coding regions and all segments of RNAs encoding proteins associated with one or more diseases, conditions or mixtures. The antisense oligonucleotides may be derived from sequences AAX55272-74. These multiple target oligonucleotides (specifically AAX55180-271) can be used for the antisense treatment of diseases and conditions. Typical diseases and conditions are those associated with impaired respiration and information in the conditions are those associated with impaired respiration and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              inflammation, including lung diseases, pulmonary vasoconstriction, inflammation, allergic rhinitis, acute asthma, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, pulmonary vasoconstriction, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukemias, lymphomas, carcinomas e.g. colon cancer, bast cancer, lung cancer, pancreatic cancer, hepatocellular carcinoma, kidney cancer, melanoma, hepatic metastases, as well as all types of cancers which may metastasize or have metastasized to the lungs, including breast and prostate cancer.
                surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pain; cystic fibrosis; allergic rhinitis;
                                                                Low adenosine antisense oligonucleotide; phosphorothicate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic;
                                                                                                                                                                                             14-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 11
                                                 respiratory obstruction; pulmonary obstruction; impeded respiration;
                                                                                                                                                            Human C/EBP
                                                                                                                                                                                                                              AAF20603;
                                                                                                                                                                                                                                                                   AAF20603 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 70; 120pp;
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17-SEP-1997;
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8; Conserv
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                                                                                                                                                        polynucleotide fragment #2170
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97US-0059160
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80.0%;
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Pred. No. 2.1e+02;
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RESULT 295
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                                                                                                                                                                                                                                                                        The present invention describes low adenosine (A) content antisense CC oligonucleotides and compositions (I) comprising them. In the antisense CC oligonucleotides the A is replaced by a 'Universal' or alternative base. CC (I) can have respiratory, bronchodilator, antiinflammatory, analyssic, immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. CC The antisense oligonucleotides and (I) can be used to down-regulate the CC expression and or activity of target polypeptides associated with CC lung/respiratory disorders and malignancies, such as stimulating and CC expression and entibodies, antibody receptors, cytokines and CC chemokines, endogenously produced specific and non-specific enzymes, in the complex proteins, adhesion molecules and their receptors, cytokine and CC chemokine receptors, adenosine receptors, bradykinin receptors, central crevous system peptide creceptors, continuing proteins, and peripheral nervous and non-nervous system peptide creceptors, brinding proteins and malignancy associated proteins. The antisense oligonucleotides may be used in this way to treat disorders including respiratory obstruction (especially pulmonary obstruction and/or bronchoconstriction) and/or lung inflammation, allergy(ies) and/or surfactant hypoproduction which are associated with a disease or condition selected from pulmonary vasoconstriction, inflammation, callergies, asthma, impeded respiration, respiratory distress syndrome (C and/or cancer. AAF18434 to AAF21543 represent human polynucleotide condition, bronchitis, and antisense oligonucleotides used in the exemplification of the present invention.
                                                                                                                                                                                                     Query Match
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Matches 8
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               AAA34481;
                                                 AAA34481 standard; DNA; 11 BP
                                                                                                                                                                                                                                                                      Sequence 11 BP; 0 A; 6 C; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Low adenosine (A) content antisense oligonucleotides which do r
trigger adenosine receptors during metabolism, useful e.g. for
cancers and respiratory obstructions -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chronic obstructive pulmonary disease; pulmonary infection; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2000-679539/66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14; Page 264; 1592pp; English.
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                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                     34.0%;
                                                                                                                                                                                                                                                                      G; 0 U; 0 other;
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                                                                                                                                                                                                                                    Score 6.8; DB 1;
                                                                                                                                                                                                                      Pred.
                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                      No. 2.1e+02;
                                                                                                                                                                                                                                   Length 11;
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                                                                                                                                                          The present invention describes a new composition comprising an antisense oligonucleotide (ON) with low adenosine (up to 15%), which CC targets nucleic acids involved in bronchoconstriction, allergies, and/or inflammation. The ON can have antiinflammatory, antiallergic, and/or can insthmatic, cycostatic and analgesic activities. The compositions are useful for the treatment of diseases associated with inflammation, comparised airways, including lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating ce.g. ischaemic conditions, pulmonary vasconstriction, allergies, cystic fibrosis, pulmonary hypertension, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukaemias, lymphomas, carcinomas, and cancers which may metastasise to the lungs, including breast and prostate cancer. The reduction of the adenosine content of the ONs reduces side effects. The A-containing ONs break down with the release of deoxyadenosine which activates adenosine receptors causing bronchoconstriction and inflammation. AAA32312 represent the curlentide sequences given in the sequence listing from the present the
                                                                          Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; adenosine receptor; low adenosine antisense oligonucleotide; phosphorothicate; impaired respiration; inflammation; allergy; allergic disease, bronchoconstriction; inhibitor; antiinflammatory; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; lung disease; ischaemic condition; pulmonary vasoconstriction; asthmatespiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; COPD; cancer; leukaemia; lymphoma; carcinoma; metastasis; ss.
                                                                                                                                                                                                         nucleotide sequences given in the sequence listing from the present invention, which correspond to SEQ ID NO:1 to 2815, and then the last 185 sequences are also called SEQ ID NO:1 to 185, but the sequences differ from the previously named sequences. SEQ ID NO:11 to 1880 (AAA332323 to AAA33992) are specifically claimed ONs from the present invention. N.B. Sequences given in the disclosure of the present invention do not match up with their corresponding SEQ ID NO: sequence
                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstriction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-AUG-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human adenosine receptor related polynucleotide SEQ ID NO:2170
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                                                                        l Similarity
8; Conserv
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ccceecccee
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                                                                                                                                                      BP;
                                                                          Conservative
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                                                                                                                                                                                               sequence listing.
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AAX55033
ID AAX550
XX AAX550
AC AAX550
XX DF O5-JUL
DT O5-JUL
DX C/EBP-
XX Antise
KW impair
KW pulmon
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ABV68762
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag
Antisense oligonucleotide; multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vasoconstriction; inflammation; allergic rhinitis;
                                                              C/EBP-beta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           e.g.
                                                                                                                          AAX55033
                                                                                                                                                                                                                                                                                                                                                     Sequence 11 BP; 1 A; 5 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 207; 1345pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-JAN-2001; 2001DE-1000127
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                                                                                                                                                    AAX55033 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Petersohn D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-DEC-2001; 2001WO-EP15179
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; skin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              fuman; skin; dermatological; vulnerary; antipsoriatic;
immunosuppressive; antiinflammatory; cytostatic; SAGE;
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8; Conserv
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                                                           antisense oligonucleotide fragment.
                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                   invention.
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                                                                                                                                                      DNA;
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Pred. No. 2.1e+02;
0; Mismatches 2
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                inflammation, including lung diseases, pulmonary vasoconstriction, inflammation, allergic rhinitis, acute asthma, allergies, asthma, inpeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, pulmonary vasoconstriction, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukemias, lymphomas, carcinomas e.g. colon cancer, breast cancer, lung cancer, pancreatic cancer, hepatocellular carcinoma, kidney cancer, melanoma, hepatic metastases, as well as all types of cancers which may metastasize or have metastasized to the lungs, including breast and prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     non-coding regions of RNAs corresponding to target genes, gene initiation codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'-end and the juxta-section between coding and non-coding regions and all segments of RNAs encoding proteins associated with one or more diseases, conditions or mixtures. The antisense oligonucleotides may be derived from sequences AAX55272-74. These multiple target oligonucleotides (specifically AAX55180-271) can be used for the antisense treatment of diseases and conditions. Typical diseases and conditions are those associated with impaired respiration and
                                                                          14-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary hypertension; pulmonary vascoonstriction; emphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; car chronic obstructive pulmonary disease; leukemia; lymphoma; car colon cancer; breast cancer; lung cancer; pancreatic cancer; colon cancer; breast cancer;
                                   Human C/EBP polynucleotide fragment #2169
                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 12
                                                                                                               AAF20602
                                                                                                                                                  AAF20602 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The specification describes antisense oligonucleotides (AAX52869-X55271) directed against at least 2 mRNAs selected from target genes, coding and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 70; 120pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides used in treatment of, e.g. pulmonary
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17-SEP-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         colon cancer; breast cancer; lung cancer; pancreatic cancer;
hepatocellular carcinoma; kidney cancer; melanoma; hepatic metastasis;
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                                                                                                                                                                                                                                                                                     3 CCGGGCCAGG
                                                                                                                                                                                                                                                                                                                       Similarity
8; Conserv
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                                                                                                                                                                                                                                                                                                                         Conservative
                                                                        (first entry)
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97US-0059160
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                                                                                                                                                  DNA;
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                                                                                                                                                                                                                                                                                                                                          Score 6.8; DB 1;
Pred. No. 2.4e+02;
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                                                                                                                                                                                                                                                                                                                                                              Length 12;
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                                                                                                                                                                                                                                                                                                                         0
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                                                                                                                                                                                                                                                                                                                         Gaps
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Low adenosine antisense oligonucleotide; phosphorothioate; allergy;

human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflamma

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The present invention describes low adenosine (A) content antisense CC oligonucleotides and compositions (I) comprising them. In the antisense CC oligonucleotides the A is replaced by a 'Universal' or alternative base. CC (I) can have respiratory, bronchodilator, antiinflammatory, analyssic, CC immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. CC The antisense oligonucleotides and (I) can be used to down-regulate the CC expression and or activity of target polypeptides associated with CC lung/respiratory disorders and malignancies, such as stimulating and CI activating peptide factors and transmitters, transcription factors, CC immunoglobulins and antibodies, antibody receptors, cytokines and CC chemokine seceptors, adenosine receptors, brinding proteins, adhesion molecules and their receptors, cytokine and CC chemokine receptors, adenosine receptors, brinding receptors, brinding proteins and paripharal nervous and non-nervous system creaptors, CNS and peripharal nervous and non-nervous system peptide constitiens eligonucleotides may be used in this way to treat disorders cincluding respiratory obstruction (especially pulmonary obstruction and/or lung inflammation, allergy(les) actimal impeded respiration, respiratory distress syndrome CC condition selected from pulmonary vasoconstriction, inflammation, colorary transplantation obstructive pulmonary disease (COPD), pain, cystic fibrosis (CP), allergic rhinitis (AR), pulmonary CD pulmonary transplantation obstructive pulmonary disease (COPD), pulmonary transplantation of contractive pulmonary disease (COPD), supertension, emphysema, chronic obstructive pulmonary disease (COPD), pulmonary transplantation of contractive pulmonary disease (COPD), pulmonary disease (COPD), allergic rhinitis (AR) pulmonary disease (COPD), pulmonary disease (COPD), contractive pulmonary disease (COPD).
     Query Match
Best Local S
Matches 8
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                                                                                                                                                                                                      Sequence 12
                                                                                                                                                                                                                                                                                                                                                   pulmonary transplantation rejection, pulmonary infections, bronchitis, and/or cancer. AAF18434 to AAF21543 represent human polynucleotide fragments and antisense oligonucleotides used in the exemplification o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 14; Page 264; 1592pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200062736-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            pulmonary hypertension; emphysema; pulmonary transplantation rejection;
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                                                         Similarity
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          Conservative
                                                                                                                                                                                                      BP; 0 A; 7 C; 5
                                                                                                                                                                                                                                                                                                    invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99US-0127958.
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80.0%;
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                                                    Score 6.8; DB 1;
Pred. No. 2.4e+02;
     Mismatches
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Query Match Best Local Similarity

34.0%; 80.0%;

Score 6.8; DB 1; Pred. No. 2.4e+02; 0 other;

Sequence

B₽;

0 A.

7 C; 5 G; 0 U;

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RESULT 299
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                                                                                     nucleotide sequences given in the sequence listing from the present invention, which correspond to SEQ ID NO:1 to 2815, and then the last 185 sequences are also called SEQ ID NO:1 to 185, but the sequences differ from the previously named sequences. SEQ ID NO:11 to 1680 (AAA32323 to AAA3392) are specifically claimed ONs from the present invention. N.B. Sequences given in the disclosure of the present invention do not match up with their corresponding SEQ ID NO: sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstriction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; adenosine receptor; low adenosine antisense oligonucleotide; phosphorothioate; impaired respiration; inflammation; allergy; allergic disease; bronchoconstriction; inhibitor; antiinflammatory; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA34480 standard; DNA; 12
                                                                                                                                                                                                                                                                                                           release of deoxyadenosine which activates adenosine receptors causing bronchoconstriction and inflammation. AAA32313 to AAA35312 represent (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 537; 1343pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-AUG-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human adenosine receptor related polynucleotide SEQ ID NO:2169
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                                                               sequence listing.
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AAX29560
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Matches 7
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                                                                                                                                                                                                           in plant biology, agriculture, human or veterinary medicine, reproduction, microbiology or environmental science, e.g. to study expression of nuclear receptors at different stages of tissue development or after treatment with particular drugs. It is also used for DNA fingerprinting (to generate products useful for differential hybridisation), or, where a 3'-anchor primer is used, to isolate the hybridisation of mRNA sequences. Sequences AAX29526-X29583 represent claimed
                                                                                                                                                                                                                                                                                                              The invention provides primers (AAX29501-X29679) for identifying sequences encoding structurally or functionally related proteins such as nuclear or G-protein coupled receptors, apoptosis-related or DNA repair/replication proteins. The identified sequences are broadly useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acid amplification; nuclear receptor; G-protein coupled receptor; apoptosis; DNA repair; DNA replication; plant biology; agriculture; human; veterinary medicine; reproduction; microbiology; hybridisation; environmental science; DNA fingerprinting; PCR primer; ss.
                                                                                                                                                                                                 primers specific for human G-protein coupled receptor genes.
                                                                                                                                                                                                                                                                                                                                                                                       Claim 21; Page 75; 92pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                    Subset of primers able to amplify group of related sequences
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Pred. No. 5.8e+02;
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         Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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US-09-371-72B-6740
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US-09-371-772B-3864
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APPLICANT: Robert A. McKay
APPLICANT: Nicholas M. Dean
ANTISENSE OLIGONUCLEOTIDE
TITLE OF INVENTION: COMPOSITONS AND METHODS FOR TH
TITLE OF INVENTION: PROTEINS
NUMBER OF SEQUENCES: 86
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Mariton
STATE: NJ
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US-08-829-954-105
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RESULT 2
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APPLICANT: Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (609) 779-2400
TELEFAX: (609) 779-8488
INFORMATION FOR SEQ ID NO: 31
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FILING DATE: August 13
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                            APPLICATION NUMBER: US/08/910,629A FILING DATE: AUGUST 13, 1997 CLASSIFICATION: 514 PRIOR APPLICATION DATA: APPLICATION NUMBER:
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COMPUTER READABLE FORM:
NEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44
MEDIUM TYPE: STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE
TITLE OF INVENTION: COMPOSITONS AND METHODS FOR THE MODULATION OF JNK
TITLE OF INVENTION: PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ANTI-SENSE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Jane Massey Licata
REGISTRATION NUMBER: 32,257
REFERENCE DOCKET NUMBER: IS
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MB
MEDIUM TYPE: STORAGE
                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 8
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TTORNEY/AGENT INFORMATION:
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                                                                                                                                                    COMPUTER: PENTIUM
OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                        STREET: 66 East
CITY: Marlton
                                                                                                                                    SOFTWARE: WORDPERFECT 6.1
                                                                                                                                                                                                                                                        COUNTRY:
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OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                             66 East Main Street
                                                                                                                                                                                                                                                           USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nicholas M. Dean
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Brett Monia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Robert A. McKay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Linear
                                                                                                                                                                                                                                                                                                                            Law Offices of Jane Massey Licata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OMBER: US/08/910,629A
August 13, 1997
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FILE REFERENCE: ISPH-0336

CURRENT APPLICATION NUMBER: US/09/209,668A

CURRENT FILING DATE: 1998-12-10

NUMBER OF SEQ ID NOS: 25

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 7

LENGTH: 20

TYPE: DNA
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            GENERAL INFORMATION:
APPLICANT: MCKAY, Robert A.
APPLICANT: MCKAY, Robert A.
APPLICANT: Dean, Nicholas M.
APPLICANT: Monia, Brett
APPLICANT: Monia, Brett
APPLICANT: Monia, Brett
APPLICANT: Nero, Pam
APPLICANT: Nero, Pam
APPLICANT: OF INVENTION: ANTISENSE OLIGONUCLEOTIDE COMPOSITIONS AND METHODS
TITLE OF INVENTION: FOR THE MODULATION OF JNK PROTEINS
TITLE OF INVENTION: FOR THE MODULATION OF JNK PROTEINS
TILLE REFERENCE: ISPH-0350
CURRENT APPLICATION NUMBER: US/09/287,796A
CURRENT FILING DATE: 1999-04-07
EARLIER APPLICATION NUMBER: 09/130,616
EARLIER FILING DATE: 1999-08-07
EARLIER FILING DATE: 1999-08-07
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Patent No. 6114517

GENERAL INFORMATION:
APPLICANT: Monia, Brett P.
APPLICANT: XU, Xiaoxing S.
TITLE OF INVENTION: METHODS OF MODULATING TUMOR NECROSIS FACTOR
TITLE OF INVENTION: alpha-INDUCED EXPRESSION OF CELL ADHESION METHODS OF MODULATING TUMOR NECROSIS FACTOR
                                                                                                                                                                                                                                                                                        Sequence 31, Application US/09287796A Patent No. 6133246
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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ARLIER APPLICATION NUMBER: 08/910,629
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les 20; Conserv
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APPLICANT: Monia, Brett
APPLICANT: Mero, Pam
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE COMPOSITIONS AND
TITLE OF INVENTION: FOR THE MODULATION OF JNK PROTEINS
FILE REFERENCE: ISPH-0350
CURRENT APPLICATION NUMBER: US/09/287,796A
CURRENT FILING DATE: 1999-04-07
EARLIER APPLICATION NUMBER: 09/130,616
FEARLIER APPLICATION NUMBER: 08/910,629
EARLIER APPLICATION NUMBER: 08/910,629
EARLIER APPLICATION NUMBER: 08/910,629
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Matches
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SEQ ID NO 42
LENGTH: 20
                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                  Sequence 31, Appli
Patent No. 6221850
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Best Local Similarity
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LENGTH: 20
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Patent No. 6133246
APPLICANT: MCKAY, Robert A.
APPLICANT: Dean, Nicholas M.
APPLICANT: Monia, Brett
APPLICANT: Monia, Brett
APPLICANT: Nero, Pam
APPLICANT: Nero, Pam
APPLICANT: Gaarde, William A.
APPLICANT: GABARDE, MISSENSE OLIGONUCLEOTIDE COMPOSITIONS AND METHODS
TITLE OF INVENTION: FOR THE MODULATION OF JNK PROTEINS
FILE REFERENCE: ISPH-0318
FULL REFERENCE: ISPH-0318
CURRENT APPLICATION NUMBER: US/09/130,616C
CURRENT FILING DATE: 1998-08-07
EARLIER APPLICATION NUMBER: 08/910,629
EARLIER FILING DATE: 1997-08-03
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                                                                                                                                                                                                                                                                                                 Sequence 6740, Application US/09371772B Patent No. 6566127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 178
SEQ ID NO 42
LENGTH: 20
                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 42, Application US/09130616C Patent No. 6221850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local
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                                  APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Rel
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/130,616C CURRENT FILING DATE: 1998-08-07 EARLIER APPLICATION NUMBER: 08/910,629 EARLIER FILING DATE: 1997-08-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JENERAL INFORMATION:
APPLICANT: MCKAY, Robert A.
APPLICANT: Dean, Nicholas M.
APPLICANT: Dean, Nicholas M.
APPLICANT: Monia, Brett
APPLICANT: Noro, Pam
APPLICANT: Nero, Pam
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE COMPOSITIONS AND METHODS
TITLE OF INVENTION: FOR THE MODULATION OF JNK PROTEINS
FILE REFERENCE: ISPH-0318
FILE REFERENCE: ISPH-0318
                                                                                                                                                                          APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
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PRIOR APPLICATION NUMBER: US 08/584,040 PRIOR FILING DATE: 1996-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
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Pred. No. 0.81;
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US-08-984-040-6244/c
US-08-984-040-6244, Application US/08584040
; Sequence 6244, Application US/08584040
; Patent No. 6346398 ;
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                                                                                                                                                       Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                COMPUTER: IM PC COMPATIBLE COMPUTER: IM PC COMPATIBLE PATENT PC-DOS/MS-DOS SOFTWARE: PATENTIN Release #1.0, Version #1.25 CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/951,923 FILING DATE: October 16, 1997 CLASSIFICATION: 435 ATTORNEY AGENT INFORMATION:

NAME: Neeley, Richard L.

REGISTRATION NUMBER: 30,092 REFERENCE/DOCKET NUMBER: BITT-001/02US

TELEPHONE: 650 843-5000

TELEPHONE: 650 843-5000
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Best Local !
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Bitter
                                                                                                                                                                                                             MOLECULE TYPE:
HYPOTHETICAL: N
ANTI-SENSE: NO
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MEDIUM TYPE: Floppy disk
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ZIP: .943
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TOPOLOGY:. lir
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                                                                                                                                                                                                                             Sequence 3004, Application US/09371772B Patent No. 6566127
                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local S
             APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (27/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 60/005,97-
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION UNMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
PRIOR APPLICATION' NUMBER: US 60/005,974
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LENGTH: 18 base pairs
TYPE: nucleic acid
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MEDIUM TYPE: 3.5" Diskette, 1.44
MEDIUM TYPE: storage
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TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM COMPUTER: IBM P.C.
OPERATING SYSTEM: IBM P.C.
OPETWARE: Word Perfect 5.1
TOTAL DATA:
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STREET: Suite 4700
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STATE: California
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633 West Fifth Street
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Stinchcomb, Dan T.
Escobedo, Jaime
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Pred. No. 5
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; TYPE: RNA
; ORGANISM: Mus sp.
US-09-371-772B-3004
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US-08-584-040-4222
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SOFTMRE: PatentIn version 3.0
SEQ ID NO 3004
LENGTH: 18
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Best Local (
                                                                                                                                                           APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/ACENT INFORMATION:
NAME: WAIDURG, Richard J.
REFERENCE/DOCKET NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1050RMATION:
TELECOMMUNICATION 1050RMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US
PRIOR FILING DATE: 1996-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dai T.
APPLICANT: Stinchcomb, Dai T.
APPLICANT: Bscobbdo, Jaine
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: CRUDITIONS RELATED TO LEVELS
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                                                                                                                                 TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: Suite 470
CITY: Los Angeles
STATE: California
                    TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                        LENGTH:
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o. 6346398
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                                                    : 17 base pairs nucleic acid
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633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Suite 4700
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                  linear
                                  single
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Pred. No. 5
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Query Match

72.0%;

Score 14.4;

BB

Length 17;

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US-09-371-772B-1989/c
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SEQ ID NO 1989
LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent No.
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions |
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 80/584,040
PRIOR FILING DATE: 1995-01-08
NUMBER OF SEQ ID NOS: 14225
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APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Stinchcomb, Dan APPLICANT: Escobedo, Jaime
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TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: RNA
ORGANISM: Homo sapiens
                                 COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
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                                                                                                                               CITY: Los Angeles
STATE: California
                                                                                                                                                                     ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
OPERATING SYSTEM:
                                                                                                           COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16 CCGGGCCAAGCCAAAG 1
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                                                                                         90071-2066
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                                                                                                                                                                                                                                                                                                                                                               McSwiggen, James
Stinchcomb, Dan T.
Escobedo, Jaime
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             IBM Compatible
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93.8%;
                                                                                                                                                                                                                                                                                                                                               METHOD AND REAGENT FOR THE
                                                                                                                                                                                                                                                                                     TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS OF VASCULAR ENDOTHELIAL
                                                                                                                                                                                                                                                                     GROWTH FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             93.8%;
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0; Mismatches
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Pred. No. 5
DOS 5.0
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RESULT 16
US-09-371-772B-4833/c
US-09-371-772B-4833/c
; Sequence 4833, Application US/09371772B
; Patent No. 6566127
; GENERAL INFORMATION: ,
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US-09-371-772B-3450/c
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                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
SEQ ID NO 3450
                                                                                                                                                                                                                Matches
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Best Local
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APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                            LENGTH: 17
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LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UNMEER: 60/005,974
FILING DATE: October 26, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
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Local Similarity 100.0%; P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
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Similarity 100.0%;
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Pred. No.
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                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                   6.9;
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; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-371-772B-5851
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US-09-371-772B-5851/c
; Sequence 5851, Application US/09371772B
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                                                                            Query Match
Best Local S
Matches 14
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Best Local S
Matches 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No. 6566127 GENERAL INFORMATION:
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LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MeBHB00, 976-J (27/37/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
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APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Rel
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/99/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-26
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APPLICANT:
APPLICANT:
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APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
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ORGANISM: Homo sapiens
  16
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                                                                              14;
                             3 CCGGGCCAGGCCAAA 17
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                                                                                                 Similarity
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                                                                                Conservative
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                                                                                                 67.0%;
93.3%;
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Pred. No. 7.
                                                                                                 Score 13.4;
Pred. No. 8;
                                                                                  Mismatches
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                                                                                                                    DB 1;
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                                                                                                                    Length 16;
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US-08-924-183-6
, Sequence 6, Application US/08924183A
, Patent No. 6218109
, GENERAL INFORMATION:
, APPLICANT: Elledge, Stephen J.

RESULT 18

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US-09-488-364-6
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                                                                                                                                                                                                                                                                                                                        RESULT 20
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US-09-488-364-6
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Matches
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CURRENT FILING DATE: 2000-01-12
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 6
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Elledge, Stephen J.
APPLICANT: Sanchez, Yolanda
TITLE OF INVENTION: MAMMALIAN CHECKPOINT GENES AND PROTEINS
FILE REFERENCE: 120541-1013
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: 120541-1003
CURRENT APPLICATION NUMBER: US/08/924,183A
CURRENT FILING DATE: 1997-09-05
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 6
LENGTH: 17
TYPE: DNA
                                                                                                                                                                                                                                                                  Sequence 1996, Application US/08584040 Patent No. 6346398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 6, Application US/09488364 Patent No. 6307015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local :
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TITLE OF INVENTION: MAMMALIAN CHECKPOINT GENES AND PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: Synthetic
                                                                                   TITLE OF INVENTION: METHOD AND REAGENT FOR THE TITLE OF INVENTION: TREATMENT OF DISEASES OR TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL TITLE OF INVENTION: GROWTH FACTOR
                                                                                                                                                                                            APPLICANT: Pavco, Pamela
APPLICANT: McSwiggen, Jav
APPLICANT: Stinchcomb, D
                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                 APPLICANT:
                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local
                                    ADDRESSEE:
                                                                                                                                                                                                                                                  INFORMATION:
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E: Lyon & Lyon
633 West Fifth Street
Suite 4700
                                                                                                                                                                                            McSwiggen, James
Stinchcomb, Dan T.
                                                                                                                                                                           Escobedo, Jaime
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93.3%; Pred. No. 8.8;
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93.3%;
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Pred. No. 8.8;
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LENGTH: 17
TYPE: RNA
ORGANISM: Homo sapiens
US-09-371-772B-541
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US-09-371-772B-541/c
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                                                                                                                                       SOFTWARE: PatentIn version SEQ ID NO 541
Best Local Similarity 93. Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 541, Application US/09371772B Patent No. 6566127
                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                       APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Rel
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pawco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
                                                                                                                                                                           NUMBER OF SEQ ID NOS: 14225
                                                                                                                                                                                          PRIOR FILING DATE: 1996-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 499-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C.
SOSTWARE: WORD Perfect 5.1
CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
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REGISTRATION NUMBER: 32
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14; Conserv
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January 11, 1996
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93.3%;
              67.0%;
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ER: 218/064
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                Score 13.4;
Pred. No. 8.
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Pred. No. 8
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Mismatches
                8.8;
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                               Length 17;
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Indels
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CCGGGCCAGGCCAAA 17

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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Disc
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Facto
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,7728
CURRENT FILING DATE: 1995-01-08
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-01-08
PRIOR FILING DATE: 1996-01-08
VUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
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US-09-371-772B-4834
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US-09-371-772B-4834/c
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LENGTH: 17
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Best Local
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                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
                    SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: McSwiggen, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 8502
                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                            ADDRESSEE: Lyon & L
STREET: 633 West Fi
STREET: Suite 4700
CITY: Los Angeles
                                                                                                                                                                                                                     STATE: California COUNTRY: U.S.A.
                                                                                    OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 7660, App
10. 6346398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION:
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Similarity 93.3%;
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633 West Fifth Street
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                                                             Word Perfect 5
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January 11, 1996
                                                                                 IBM P.C.
                    US/08/584,040
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Pred. No. 8.8;
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PRIOR FILING DATE: 1996-01-08
; NUMBER OF SEQ ID NOS: 14225
; SOFTWARE: Patentin version 3.0
; SEQ ID NO 3449
; LENGTH: 17
; TYPE: RNA
; ORGANISM: Mus Sp.
US-09-371-772B-3449
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                                                                              US-08-584-040-5795/c
; Sequence 5795, Application US/08584040
; Patent No. 6346398
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TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 76
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                      Query Match
Best Local (
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 08/584,040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -09-371-772B-3449/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: WARDURG, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17 CCAGGCCAAAGTC 5
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Pavco, Pamela
McSwiggen, James
Stinchcomb, Dan
                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                      65.0%; Score 13; 100.0%; Pred. No.
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Re

APPLICANT: APPLICANT: APPLICANT:

Escobedo,

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RESULT 26
US-09-371-772B-2661/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local :
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-U (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1995-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 514

PRIOR APPLICATION DATA:
RAPPLICATION UNMBER: 60/005,974

FILING DATE: October 26, 1995

ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.

REGISTRATION NUMBER: 32,327
                                                                                                                                                                                       APPLICANT: Ribozyme Pharmaceuticals, Inc
APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
                                                                                                                                                                                                                                                                                              ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: torrage
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/584,040 FILING DATE: January 11, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                           16
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                                                                                                                                                                               MCSwiys
Stinchcomb, Dan
Stinchcomb, Jaime
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              64.0%;
87.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TREATMENT OF DISEASES OR CONDITIONS RELATED TO LE OF VASCULAR ENDOTHELIAL
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Pred. No. 1:
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REFERENCE/DOCKET NUMBER: 39,911

REFERENCE/DOCKET NUMBER: 96700/437

TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 697-5995
TELEX: (212) 286-0854 or 286-0082
TELEX: TWX 710-581-4766
INFORMATION FOR SEQ ID NO: 49:
SEQUENCE CHARACTERISTICS:
LENGTH: 12
TYDE:
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; ORGANISM: Mus sp.
US-09-371-772B-2661
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                                                                    Best
                                                          Matches
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                                                                                     Query Match
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/0
FILING DATE: MARCH 20, 1
ATTORNEY/AGENT INFORMATION:
                                                                                                                                MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: WIILIAM R. JACOBS, JR., JAMES M. MUSSER AND APPLICANT: AMALIO TELENTI
                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: 3.5 INCH 1.44 N
MEDIUM TYPE: DISKETTE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: MS-DOS
SOFTWARE: ASCII
                                                                      Local
                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                         STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                          NAME: ELIZABETH A. BOGOSIAN REGISTRATION NUMBER: 39,911
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 90 PARK AVENUE CITY: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
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No. 6015890
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                             5 GGGCCAGGCCAA 16
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                                                                      Similarity
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                                                                                                                                                                                          nucleic acid
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                                                          Conservative
                                                                                                                                                             linear
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                                                                                                                              other nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              64.0%;
                                                                      52.0%; Score 10.4;
91.7%; Pred. No. 1
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Pred. No. 1:
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                                                                                     DB 1; Length 12;
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                                                          Indels
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                                                      Gaps
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US-09-281-418-100 ; Sequence 100, Application US/09281418 ; Patent No. 6287769

RESULT 28

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US-08-882-046-77
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 100
LENGTH: 12
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                                                                                                                                                                        APPLICATION NUMBER: US/08/882,04
FILING DATE: 25-UN-1997
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-UW 26
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE: 9982-7
CURRENT APPLICATION NUMBER: US/09/281,418
CURRENT FILING DATE: 1999-03-30
EARLIER APPLICATION NUMBER: JP/1998/87651
EARLIER FILING DATE: 1998-03-31
EARLIER FILING DATE: 1998-03-16
EARLIER FILING DATE: 1999-03-16
                                                                                            TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Primer
                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 THE OF INVENTION: Method of Amplifying DNA Fragment, Appar TTLE OF INVENTION: agment, Method of Assaying Microorganism TTLE OF INVENTION: nisms and Method of Assaying Contaminant
                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Rel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Spinner, Nancy B.
TITLE OF INVENTION: Human Jagged Polypeptide, Encoding
TITLE OF INVENTION: Nucleic Acids and Methods of Use
NUMBER OF SEQUENCES: 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Campbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Suite
CITY: San Diego
STATE: California
                                                     TYPE: nucleic acid
                                     STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
                                                                                                                                                          TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    9 CAGGCCAAAGTC 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NOS: 216
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Krantz, Ian D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Li, Linheng
                    linear
                                                                                                                                                          (619) 535-9001
                                                                                                                                                                                                                                                                                                UMBER: US/08/882,046
25-JUN-1997
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91.7%;
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Pred. No. 17
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                                                                                                             ; ORGANISM: synthetic US-09-922-445-36
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APPLICANT: Andersson, M
APPLICANT: Berglund, L
APPLICANT: Reneland, R
APPLICANT: Adam, Gail
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-922-445-11
                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/922,445
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 51
SOFTWARE: PatentIn version 3.1
SEQ ID NO 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/922,445
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 51
SOFTWARE: PatentIn version 3.1
SEQ ID NO 11
LENGTH: 13
TYPE: DNA
                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                           Sequence 36, Application US/09922445 Patent No. 6528268
                                                      Matches
                                                                                 Query Match
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Best Local Similarity
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Patent No. 6528268
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                                                                                                                                                                                                                                          TITLE OF INVENTION: REAGENTS AND METHODS FOR DETECTION OF HEART FAILURE FILE REFERENCE: GG126US
                                                                                                                                                                                                                                                                     APPLICANT: Andersson,
APPLICANT: Berglund,
APPLICANT: Reneland,
APPLICANT: Adam, Gail
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: GG126US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: REAGENTS AND METHODS FOR DETECTION OF HEART FAILURE
                                                                                                                                        LENGTH: 13
TYPE: DNA
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                          3 CCGGGCCAGGCCA 15
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                                                                    Similarity
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                                                      Conservative
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                                                                                                                                                                                                                                                                                 , Lars G. T.
, Rikard H.
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Rikard H.
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                                                                    Score 9.8; I
Pred. No. 23,
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Pred. No. 23
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Pred. No. 23;
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US-09-49-155A-324 ; Sequence 324, Application US/09249155A ; Patent No. 6538173

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SEQ ID NO 324
LENGTH: 11
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Best Local Similarity
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PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: US 60/097,937
PRIOR FILING DATE: 1998-08-26
PRIOR APPLICATION NUMBER: US 60/102,051
PRIOR FILING DATE: 1998-09-28
PRIOR FILING DATE: 1998-09-28
NUMBER OF SEQ ID NOS: 346
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TITLE OF INVENTION:
FILE REFERENCE: 004
                                                                                                                                   TELEPHONE: 650 843-500
TELEFAX: 650 857-0663
TELEX: 380816COOLEYPA
INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                    NAME: Neeley, Richard L.
REGISTRATION NUMBER: 30,092
REFERENCE/DOCKET NUMBER: BIT
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650 843-5000
                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Bitter, TITLE OF INVENTION:
                                                                                                                     SEQUENCE CHARACTERISTICS:
              HYPOTHETICAL:
                                MOLECULE TYPE:
                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 5 L. CITY: Palo Alto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
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                                         TYPE:
STRANDEDNESS: BIT
TOTOGY: linear
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                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/951,923
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STREET: 5 Palo Alto Square, 3:
                                                                                                     ENGTH:
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                                                                                 nucleic acid
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Pred. No. 22;
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                                                                             US-08-488-551B-387
                                                                                         RESULT 35
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                                                 Sequence 387, Application US/08488551B Patent No. 6015661
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                                     GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US/08/388
FILING DATE: 14-FEB-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: DiGLIGIO, Frank S.
REGISTRATION NUMBER: 31,346
REFERENCE/DOCKET NUMBER: 9606
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                STRANDEDNESS: single TOPOLOGY: linear MOLECULE TYPE: DNA (general contents)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (516) 742-4366
                                                                                                                                                                                                 Local
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ZIP: 11530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Garden City
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400 Garden City Plaza
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Crowe, Suzanne
             Dale A. McPhee
                          Nicholas J. Deacon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      United States
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83.3%;
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Pred. No. :
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Pred. No. 43;
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NON-PATHOGENIC STRAINS OF HIV-1

TUMBER OF SEQUENCES:

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Query Match
Best Local Similarity
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US-08-522-384-63/c
                                                                               SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 63
                                                                                                                                                                                                                                                             Sequence 63, Application US/08522384 Patent No. 6110667
                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                               TITLE OF INVENTION: PROCESSES, APPARATUS AND COMPOSITIONS FOR TITLE OF INVENTION: CHARACTERIZING NUCLEOTIDE SEQUENCES FILE REFERENCE: 2458-4029
                                                                                                                CURRENT FILING DATE: 1996-11-15
NUMBER OF SEQ ID NOS: 122
                                                                                                                                                CURRENT APPLICATION NUMBER: US/08/522,384
                                                                                                                                                                                                               APPLICANT: LOPEZ-NIETO, CARLOS APPLICANT: NIGAM, SANJAY KUMAR
                    TYPE: DNA
ORGANISM: Unknown Organism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (516) 742-431
OTHER INFORMATION: Description of Unknown Organism:
                  FEATURE:
                                                                  ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS LENGTH: 10 base pairs
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4343
TELEFAX: (516) 742-4366
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: FRANK S. DIGIGLIO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Release #1.0, Version #1.25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: NEW YORK COUNTRY: U.S.A. ZIP: 11530-0299
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/488,551B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: GARDEN CITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
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400 GARDEN CITY PLAZA
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90.0%;
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Pred. No. 28;
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                                                                                                                                                                                                                                                                                                                                                                                                   US-08-822-586-49
                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: POSEQ ID NO 24
                                                                                                                                                                                                                                                                                                                                                                       Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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Best Local
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APPLICANT: KOHARA, MICHINORI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: A VECTOR EXPRESSING AN RNA VIRAL FULL-LENGTH GENE AND TITLE OF INVENTION: ITS USE FILE REFERENCE: 04853.0051-0000 CURRENT APPLICATION NUMBER: US/09/720,201A CURRENT FILING DATE: 2000-12-22 PRIOR APPLICATION NUMBER: US/012-22 PRIOR APPLICATION NUMBER: US/012-22 PRIOR APPLICATION NUMBER: US/012-22 PRIOR FILING DATE: 1998-06-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-720-201A-24
                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 10
TYPE: DNA
ORGANISM: Artificial Sequence
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NUMBER OF SEQ ID NOS: 25
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CURRENT AFFLICATION DATA:
APPLICATION NUMBER: US/08/822,586
FILING DATE: MARCH 20, 1997
                                                                                                  ZIP: 10016
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 INCH 1.44 Mb STORAGE
MEDIUM TYPE: DISKETTE
                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 5
                                                                                                                                                                                                                                                                                     TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                    COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: MS-DOS SOFTWARE: ASCII
                                                                                                                                                                                      STREET: 90 PARI
CITY: NEW YORK
STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                     COUNTRY: U.S.A.
                                                                                                                                                                                                                                     ADDRESSEE:
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                                                                                                                                                                                    NEW YORK
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TAIRA, KAZUNARI
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                                                                                                                                                                                                                                                                                                                                                                                      Application US/08822586
                                                                                                                                                                                                                    90 PARK AVENUE
                                                                                                                                                                                                                                                                                                                       AMALIO TELENTI
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                                                                                                                                                                                                                                                                                     MUTANTS THEREOF
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90.0%;
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90.0%;
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Pred. No. 28
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Pred. No. 28;
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US-08-822-586-49
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 697-5995
TELEFAX: (212) 286-0854 or 286-0082
TELEFAX: TWX 710-581-4766
INFORMATION FOR SEQ ID NO: 49:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Paul, Thomas D.
REGISTRATION NUMBER: 32,714
REFERENCE/DOCKET NUMBER: D-5900
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713/651-5325
TELEPHONE: 713/651-5326
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                                                                                                                                   TELEFAX: 713/651-5246
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tent No. 6083695
                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPOTER:
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/859,954
APPLICATION NUMBER: US/08/859,954
                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: ot
HYPOTHETICAL: NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Hardin, Susan H. APPLICANT: Homayouni, Ramin APPLICANT: Hardin, Paul E.
                                       MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE
                                                                        STRANDEDNESS:
                                                                                            TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U.S.A.
ZIP: 77010-3095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Houston
STATE: Texas
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: Fulbright & Jaworski L.L.P.
1301 McKinney, Suite 5100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Application US/08859954
                                                         linear
other nucleic acid
/desc = "oligonucleotide"
YES
                                                                          single
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Pred. No. 34;
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Query Match
Best Local Similarity
Thes 8; Conserv:
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Best Local Similarity luu
Conservative
                                                                                                                                    TELEPAX: 650-326-2422
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 183,
Patent No. 58
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: Floppy disk
COMPUTER: PA PC COMPATIBLE
COMPUTER: PC COMPATIBLE
COMPUTER: PC COMPATIBLE
COMPATIBLE
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/441,887A
FILING DATE: 16-MAY-1995
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
                                                                                                                                                                                                                                            NAME: Liebeschuetz, Joseph O.
REGISTRATION NUMBER: 37,505
REFERENCE/DOCKET NUMBER: 018547-004160US
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-326-2400
                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/143,312
FILING DATE: 26-OCT-1993
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/082,937
APPLICATION NUMBER: US 08/082,937
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, 8th Floor
                                                                                                                                                                                                                                                                                                                                      FILING DATE: 25-JUN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Sheldon, Edward L.
TITLE OF INVENTION: Arrays of Nucleic Acid Probes on
TITLE OF INVENTION: Biological Chips
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: San Francisco
STATE: California
                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY:
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    Application US/08441887A
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                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cronin,
                                                                                                                           linear
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Huang, Xiaohua X.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chee, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lipshutz,
                                                                                                       DNA (probe)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Maureen T
              40.0%; Score 8; 1
100.0%; Pred. No.
tive 0; Mismatc
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Pred. No. 1.6e+02;
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                  Mismatches
                                    DB 1;
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                                                   Length 10;
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                                                                                                                                                           Patent No. 6010895
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 19
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                       APPLICANT:
                                                                                                                     APPLICANT:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
                           CORRESPONDENCE ADDRESS:
                                                                                                                                 APPLICANT: Deacon, Nicholas J
                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
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CITY: Los Angeles
                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic
STRANDEDNESS:
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OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Warburg, Richard REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE:
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                ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                  nucleic acid
                                                          INVENTION: NON-PATHOGENIC STRAINS OF HIV-1
                                                                                    Learmont, Jennifer C. McPhee, Dale A. Crowe, Suzanne
3: Scully, Scott, Murphy & Presser
400 Garden City Plaza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (213) 955-0440
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                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SYSTEM: IBM P.C. DOS (Version 5.0)
WordPerfect (Version 5.1)
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100.0%; Pr
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METHOD AND REAGENT FOR
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Pred. No.
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Best Local Similarity
Watches 8; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3
Patent No.
APPLICATION NUMBER: US/08/388
FILING DATE: 14-FEB-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Dic3glio, Frank 5.
REGISTRATION NUMBER: 31,346
REFERENCE/DOCKET NUMBER: 9606
TELECOMMUNICATION: INFORMATION:
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                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/388,353
FILING DATE: 14-FBB-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Digiglio, Frank S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 10 base pairs
                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION: NON-PATHOGENIC STRAINS OF HIV-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 96
TELECOMMUNICATION INFORMATION:
TELEPHONE: (516) 742-4343
TELEPHAX: (516) 742-4366
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                                                                                                                                                                                                                                                                                                       : Garden City
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400 Garden City Plaza
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McPhee, Dale A.
                                                                                                                                                                                                                                                                   United States
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; TOPOLOGY: 1; MOLECULE TYPE: US-08-488-551B-388
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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                        APPLICATION NUMBER: PM4002
FILING DATE: 21-FEB-1994
APPLICATION NUMBER: PN0284
FILING DATE: 23-DEC-1994
APPLICATION NUMBER: US 08/
FILING DATE: 11-FEB-1995
APPLICATION NUMBER: PN3021
APPLICATION NUMBER: PN3021
APPLICATION NUMBER: PN3021
APPLICATION NUMBER: PN3021
APPLICATION STORMATION:
NAME: FRANK S. DIGIGLIO
REFERENCE/DOCKET NUMBER: 9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                equence 388, Application US/08488551B
                                                                                                                                                                          TELEFAX: (516) 742-4366 INFORMATION FOR SEQ ID NO:
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                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,551B
FILING DATE: 07-JUN-1995
PRICE APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy
                                                                                                                                                                                                         TELEPHONE: (516) 742-4343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
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hes 8; Conservative (
                                                                                                                    TYPE: nucleic acid
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APPLICATION NUMBER: PM3807 ....

FILING DATE: 14-FEB-1994

FILING DATE: 14-FEB-1994

AND DATE: PM4002 (AU)
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OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                     STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U.S.A. ZIP: 11530-0299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 10 base pairs TYPE: nucleic acid STRANDEDNESS: single TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E: SCULLY, SCOTT, MURPHY & PRESSER 400 GARDEN CITY PLAZA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              David Cooper
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nicholas J. Deacon
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40.0%; 5
100.0%;
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Score 8; 1
Pred. No.
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Pred. No.
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DB 1;
5. 32;
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o. 32;
              Length 10;
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                            US-08-924-183-6/c
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Sequence 6, Application US/08924183A Patent No. 6218109
                                                                                                                                                     Matches
                                                                                                                                                                Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                        TELEFAX: (516) 742-43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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APPLICATION NUMBER:
FILING DATE: 14-FEB-
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MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
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NAME: FRANK S. DIGIGLIO
                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 23-DEC-1994
APPLICATION NUMBER: US 08,
FILING DATE: 14-FEB-1995
APPLICATION NUMBER: PN3021
FILING DATE: 17-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: NON-PATHOGENIC STRAINS OF HIV-1
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                                                                                                                                                                                                                                                                     TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 11530-0299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: GARDEN CITY
                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 21-FEI APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
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                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                     (516) 742-4366
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dale A. McPhee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nicholas J. Deacon
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                                                                                                                                                                                                                                                                                                                                                   (516) 742-4343
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     14-FEB-1994
UMBER: PM4002 (AU)
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                                                                                                                                                Score 8; DB 1; Pred. No. 32; 0; Mismatches
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APPLICANT: Elledge, Stephen J.

COMPUTER READABLE FORM:

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RESULT 48
US-08-910-629A-31/c
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Matches
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 6
LENGTH: 17
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CURRENT FILING DATE: 2000-01-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Elledge, Stephen J.
APPLICANT: Sanchez, Yolanda
TITLE OF INVENTION: MAMMALIAN CHECKPOINT GENES AND PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/08/924,183A
CURRENT FILING DATE: 1997-09-05
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILE REFERENCE: 120541-1013
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TITLE OF INVENTION: MAMMALIAN CHECKPOINT GENES AND PROTEINS
                                                                                         TITLE OF INVENTION: ANT
TITLE OF INVENTION: COM
TITLE OF INVENTION: PRO
NUMBER OF SEQUENCES: 86
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
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ORGANISM: Artificial Sequence
FEATURE:
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                                                           STREET:
                  COUNTRY:
                                                                              ADDRESSEE:
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08053
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                                                             66 East Main Street
                USA
                                                                                                                                                                                          Nicholas M. Dean
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                                                                                                                                                                         Brett Monia
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                                                                                                                                                                                                          Robert A. McKay
                                                                            Law Offices of Jane Massey Licata
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81.8%;
                                                                                                                          ANTISENSE OLIGONUCLEOTIDE COMPOSITONS AND METHODS FOR THE MODULATION OF JNK PROTEINS
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81.8%;
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Pred. No. !
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; ANTI-SENSE:
US-08-910-629A-31
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US-08-910-629A-42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No.
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Best Local
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              FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Jane Massey Licata
                                                         CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE,
MEDIUM TYPE: STORAGE
                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                      SOFTWARE: WORDPERFECT 6.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
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PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: WORDPERFECT CURRENT APPLICATION DATA
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: Nucleic Acid
STRANDEDNESS: Sing
NAME: Jane Massey Licata
REGISTRATION NUMBER: 32,
                                                                                         APPLICATION NUMBER: US/08/91 FILING DATE: August 13, 1997 CLASSIFICATION: 514
                                                                                                                                                               OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                           CITY: .Marlton
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OPERATING SYSTEM: WINDOWS 95
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REGISTRATION NUMBER: 32,
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10; Conserv
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5877309
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66 East Main Street
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Brett Monia
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                                                                                                                                                                                                                                                                                                                                                                                                      ANTISENSE OLIGONUCLEOTIDE
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 32,257
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Pred. No. 55;
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; ANTI-SENSE:
US-08-910-629A-42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 7
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 7, Application US/09209668A Patent No. 6114517
                                                                                                                                                                                                                                                                                         Sequence 31, Application US/09287796A Patent No. 6133246
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APPLICANT: Nero, Pam
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE COMPOSITIONS AND METHODS
TITLE OF INVENTION: FOR THE MODULATION OF JNK PROTEINS
TILE REFERENCE: ISPN-0350
CURRENT APPLICATION NUMBER: US/09/287,796A
CURRENT FILING DATE: 1999-04-07
EARLIER APPLICATION NUMBER: 09/130,616
EARLIER APPLICATION NUMBER: 08/910,629
EARLIER FILING DATE: 1997-08-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Monia, Brett P.
APPLICANT: Xu, Xiaoxing S.
TITLE OF INVENTION: METHODS OF MODULATING TUMOR NECROSIS FACTOR
TITLE OF INVENTION: METHODS OF MODULATING TUMOR NECROSIS FACTOR
TITLE OF INVENTION: alpha-INDUCED EXPRESSION OF CELL ADHESION MO
FILE REFERENCE: ISPH-0336
CURRENT FELLORITON NUMBER: US/09/209,668A
CURRENT FILING DATE: 1998-12-10
NUMBER OF SEQ ID NOS: 25
                                                                                                                                                                                       SENERAL INFORMATION:
APPLICANT: MCKAY, ROBERT A.
APPLICANT: Dean, Nicholas M.
APPLICANT: Monia, Brett
APPLICANT: Mero, Pam
APPLICANT: Mero, Pam
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STRANDEDNESS: Sing:
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10; Conservative
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0; Mismatches
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Pred. No. 55;
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US-09-130-616-31/c
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LENGTH: 20
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Best Local Similarity
CURRENT APPLICATION NUMBER: US/09/130,616C CURRENT FILING DATE: 1998-08-07 EARLIER APPLICATION NUMBER: 08/910,629 EARLIER FILING DATE: 1997-08-03 NUMBER: OF SEQ ID NOS: 178
                                                                                                                                                                            APPLICANT: MCKAY, Robert A.
APPLICANT: Dean, Nicholas M.
APPLICANT: Monia, Brett
APPLICANT: Nero, Pam
APPLICANT: Nero, Pam
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE COMPOSITIONS
TITLE OF INVENTION: FOR THE MODULATION OF JNK PROTEINS
FILE REFERENCE: ISPH-0318
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APPLICANT: Nero, Pam
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE COMPOSITIONS
TITLE OF INVENTION: FOR THE MODULATION OF JNK PROTEINS
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TYPE: DNA
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Pred. No. 55;
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LENGTH: 20
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GENERAL INFORMATION:
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CURRENT FILING DATE: 1998-08-07
EARLIER APPLICATION NUMBER: 08/910,629
EARLIER FILING DATE: 1997-08-03
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APPLICANT: Mean, Nicholas M.
APPLICANT: Monia, Brett
APPLICANT: Nero, Pam
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: ANTISENSE OLIGONUCLECTIDE COMPOSITIONS AND METHODS
TITLE OF INVENTION: FOR THE MODULATION OF JNK PROTEINS
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ORGANISM: Artificial Sequence
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           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                          NUMBER OF SEQUENCES: 4
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                                                                                       COUNTRY:
                                                                                                             CITY: New York
STATE: NY
                                                                                                                                            STREET:
                                                                                                                                                         ADDRESSEE: Courtis, Morris & Safford ADDRESSEE: c/o William S. Frommer
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Pred. No. 55;
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Best Local Similarity
7; Conserva
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US-08-709-209-423
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 423, Application US/08709209 Patent No. 5762938
                                                                    CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 08/105,483
FILING DATE: 12-AUG-1993
APPLICATION NUMBER: US 07/847,951
FILING DATE: 06-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-240
THEFENNEY. 2172) 840-2323
THEFENNEY. 2172) 840-2323
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TELEPHONE: (212) 840-33:
TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
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PRIOR APPLICATION UMBER: US 07/847,951
APPLICATION SOME OF THE OFFICE OF T
                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 530 Fif
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/709,209 FILING DATE: 21-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Courtis, Morris & Safford ADDRESSEE: C/O William S. Frommer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 8 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Frommer, William S. REGISTRATION NUMBER: 25,5 REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/105,483 FILING DATE: 12-AUG-1993
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                                                            840-3333
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100.0%; Pred. No.
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                                                                                                  454310-2400
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RESULT 58
US-08-458-101-423
; Sequence 423, Application US/08458101
; Patent No. 5766599
; GENERAL INFORMATION:
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US-08-303-275-131
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TELEPHONE: (212) 840-333
TELEPEAX: (212) 840-0712
INFORMATION FOR SEQ ID NO: 131:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 7; Conserv
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Best Local Similarity 100.0%;
Matches 7; Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/303,275
FILING DATE:
CLASSIFICATION: 424
ERIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/897,382
FILING DATE: 11-JUN-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Paolet
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: IMMUNODEFICIENCY VIRUS RECOMBINANT TITLE OF INVENTION: POXVIRUS VACCINE
                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 45
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CITY: New York
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TOPOLOGY: li
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Curtis, Morris & Safford
ADDRESSEE: c/o William S. Frommer
                                                                                                                                                              11 GGCCAAA 17
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100.0%; Pr
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; Pred. No.
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o. 1.6e+02;
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US-08-859-954-106
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                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Hardin, Susan H.
APPLICANT: Homeyouni, Ramin
APPLICANT: Hardin, Paul E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (212) 840-33.
TELEFAX: (212) 840-0712
INFORMATION FOR SEQ ID NO:
                                                                     APPLICANT: Hardin, Paul E.
TITLE OF INVENTION: Design
TITLE OF INVENTION: Gene
SOUTH OF SEQUENCES: 566
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: 550
CITY: New York
CTATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/00 FILING DATE: 01-JUN-1995 CLASSIFICATION: 424 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS
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                                         STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Frommer, William S.
REGISTRATION NUMBER: 25,506
REFERENCE/DOCKET NUMBER: 454310-2740
                                                         ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
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ZIP: 10036
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Curtis, Morris & Saffo
ADDRESSEE: c/o William S. Frommer
STREET: 530 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 8 base pairs
                                                                                                                                                                                                                                                                                                                                           11 GGCCAAA 17
        Houston: Texas
                                                                                                                                                                                                                                                                                                               GGCCAAA 7
                                       1301 McKinney, Suite 5100
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Pincus, Steven E.
Cox, William I.
Audonnet, Jean-Christophe Francis
Gettig, Russell Robert
VENTION: GENETICALLY ENGINEERED VACCINE
VENTION: STRAIN
                                                                                                                                                                                                                                                                                                                                                                                Conservative
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de Taisne, Charles
Limbach, Keith J.
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                                                           Fulbright & Jaworski L.L.P
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                                                                                        Design and Optimized Primer Library
Gene Sequencing and Method Thereof
566
                                                                                                                                                                                                                                                                                                                                                                                               35.0%;
100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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Gaps

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Query Match
Best Local Similarity
Thes 7; Conserve
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US-08-859-954-106
                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
APPLICANT:
                            FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/859,954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "oligonucleotide"
HYPOTHETICAL: YES
ANTI-SENSE: YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Design and Optimized Primer Library for TITLE OF INVENTION: Gene Sequencing and Method Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION: TELEPHONE: 713/651-5325
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                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES:
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PRIOR APPLICATION DATA:
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TTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/859,954 FILING DATE:
                                                                                                                                                                                                              COUNTRY: U.S.A. ZIP: 77010-3095
                                                                                                                                                                                                                                         STATE: Texas
                                                                                                                                                                                                                                                              CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 8 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                             STREET:
                                                                                                                                                                                                                                                                                              ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12 GCCAAAG 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 GCCAAAG 7
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77010-3095
                                                                                                                                                                                                                                                                                                                                                                                                                                                6083695
                                                                                                                                                                                                                                                                           3: Fulbright & Jaworski L.L.P. 1301 McKinney, Suite 5100
                                                                                                                                                                                                                                                                                                                                                                                                            Hardin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Application US/08859954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                          Hardin, Paul E.
                                                                                                                                                                                                                                                                                                                                                                                           Homayouni, Ramin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         35.0%;
                                                                                                                                                                                                                                                                                                                            566
                                08/632,782
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 7;
Pred. No.
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Mo. 1.6e+02;
0;
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TELEFAX: 713/651-5246
INFORMATION FOR SEQ ID NO:
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                                                                                                    INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DESCRI. YES
                                                                                                                                                 NAME: Fraeer, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
                                                                                                                                                                                                                                                                       FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/240,382
MOLECULE TYPE:
                                                                                SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/2/
FILING DATE: 10 May 1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PS/2 Model 50Z or 55S
OPERATING SYSTEM: MS_DOS (Version 5
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 45
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "oligonucleotide"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: 3.5" DISKELLE, - S5SX COMPUTER: IBM PS/2 Model 50Z or 55SX COMPUTER: MS-DOS (Version 5.1)
                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
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                                                                                                                                   TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 8 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER:
            TOPOLOGY:
                                                              LENGTH:
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                                              nucleic acid
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                                                              8 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Wakita, Takaji
                                                                                                                                    (617) 542-8906
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%; Pred. No. 1.6
0; Mismatches
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Query Match
Best Local Similarity
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                                                               RESULT 63
US-08-859-954-107
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US-08-859-954-56
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Sequence 107, Application US/08859954
Patent No. 6083695
GENERAL INFORMATION:
APPLICANT: Hardin, Susan H.
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Best Local Similarity
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NAME: Paul, Thomas D.
REGISTRATION NUMBER: 32,714
REFERENCE/DOCKET NUMBER: D-59
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713/651-5325
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                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 713/651-5246
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                        MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "oligonucleotide" HYPOTHETICAL: YES ANTI-SENSE: YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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APPLICANT: Hardin, Paul E.
TITLE OF INVENTION: Design and Optimized Primer Librar
TITLE OF INVENTION: Gene Sequencing and Method Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
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TTY: Houston
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                                                                                                                                                                                                                                                                                                                                                     LENGTH: 8 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
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ZIP: 77010-3095
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60836
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                                                                                                                               GGCAAAGT 8
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1301 McKinney, Suite 5100
                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                           linear
                                                                                                                                                                                                             32.0%;
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                                                                                                                                                                                            Score 6.4; DB 1; Length 8; Pred. No. 1.6e+02; 0; Mismatches 1; Indels
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Earn, Scott
APPLICANT: Dai, Jia Le
APPLICANT: Dai, Jia Le
APPLICANT: Vogelstein, Bert
APPLICANT: Vogelstein, Bert
APPLICANT: Vogelstein, Bert
APPLICANT: Vogelstein, Bert
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Human SMAD3 and SMAD4 are
TITLE OF INVENTION: Sequence-specific transcription activators
FILE REFERENCE: 01107.74098
CURRENT APPLICATION NUMBER: US/09/041,675A
CURRENT FILING DATE: 1998-03-13
NUMBER OF SEQ ID NOS: 27
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Watches 7; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                      US-09-041-675-9/c
                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 64
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                                                                                                                                                                                                                                                                                                                                                          Sequence 9, Application US/09041675A Patent No. 6100032
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DESCRIPTION YES
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Hardin, Paul E.
TITLE OF INVENTION: Design and Optimized Primer Library for TITLE OF INVENTION: Gene Sequencing and Method Thereof NUMBER OF SEQUENCES: 566
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fulbright & Jaworski L.L.P.
STREET: 1301 McKinney, Suite 5100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 8 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
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APPLICATION NUMBER: 08/6:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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REGISTRATION NUMBER: 32,714
REFERENCE/DOCKET NUMBER: D-5900
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       YES
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Pred. No. 1.6e+02;
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APPLICANT: Vogelstein, Bert
APPLICANT: Winzler, Kenneth
TITLE OF INVENTION: Human SMAD3 and SMAD4 are
TITLE OF INVENTION: Human SMAD3 and SMAD4 are
TITLE OF INVENTION: sequence-specific transcription activators
FILE REFERENCE: 01107.74098
CURRENT APPLICATION NUMBER: US/09/041,675A
CURRENT FILING DATE: 1998-03-13
NUMBER OF SEQ ID NOS: 27
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 15
                                                                                                       SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 23
                                                                                                                                                                                                                                                                                                                                                  Sequence 23, Application US/09041675A Patent No. 6100032
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Best Local
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Best Local
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APPLICANT: Zawel, Leigh
APPLICANT: Zawel, Leigh
APPLICANT: Dai, Jia Le
APPLICANT: Wogelstein, Bert
APPLICANT: Kinzler, Kenneth
TITLE OF INVENTION: Human SMAD3 and SMAD4 are
TITLE OF INVENTION: sequence-specific transcription activators
FILE REFERENCE: 01107.74098
                                                                                                                                                                                                                                                                                                                                                                                    -09-041-675-23
                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/041,675A
                                                                                                                                         NUMBER OF SEQ ID NOS:
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APPLICANT: Zawel, Leigh
                                       LENGTH: 8
TYPE: DNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
OTHER INFORMATION: synthetic random oligonucleotides selected for OTHER INFORMATION: binding to human SMAD3 or human SMAD4
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Pred. No. 1.6e
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human SMAD3 or human SMAD4
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Pred. No. 1
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.6e+02;
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CURRENT APPLICATION NUMBER: US/08/646,301A; CURRENT FILING DATE: 1996-05-16; NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 10
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TITLE OF INVENTION:
TITLE OF INVENTION:
FILE REFERENCE: PB15
Query Match
Best Local S
Matches 4
                                                                                                                                                                                    FILE REFERENCE: PB1508USW
CURRENT APPLICATION NUMBER: US/08/646,301A
CURRENT FILING DATE: 1996-05-16
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Patentin Ver: 2.1
SEQ ID NO 10
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APPLICANT: Huber, Brian E.
TITLE OF INVENTION: Transc
Patent No. 6194211 Antige
TITLE OF INVENTION: Antige
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Matches
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OTHER INFORMATION: sequence
Patent No. 6194211
                                                                              Patent No.
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Best Local
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                                                                         OTHER INFORMATION: Description of Artificial Sequence: consensus OTHER INFORMATION: sequence B4 from DNA Sequence 1:3-11 (1990).atent No. 6194211
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TYPE: DNA
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APPLICANT: Huber, Brian E.
                                                                                                                                       LENGTH: 8
TYPE: DNA
ORGANISM: Artificial Sequence
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              32.0%;
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Score 6.4; DB Pred. No. 1.6e-4; Mismatches
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Pred. No. 1.6e+02;
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Pred. No. 1
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             DB 1;
1.6e+02;
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                                              RESULT 71
US-08-522-384-63
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US-09-398-499-16/c
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Best Local S
Matches 7
Sequence 63, Application US/08522384 Patent No. 6110667 GENERAL INFORMATION:
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Best Local Similarity
Matches 7; Conserv
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LENGTH: 8
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CURRENT APPLICATION NUMBER: US/09/398,499
CURRENT FILING DATE: 1999-09-17
PRIOR APPLICATION NUMBER: 60/101,011
PRIOR FILING DATE: 1998-09-18
NUMBER OF SEQ ID NOS: 58
                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Benson, Andrew K.
TITLE OF INVENTION: HIGH RESOLUTION GENOY
FILE REFERENCE: UNL 293
CURRENT APPLICATION NUMBER: US/09/398,499
CURRENT FILING DATE: 1999-09-17
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acent No. 6284466
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PRIOR FILING DATE: 1998-09-18
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TITLE OF INVENTION: HIGH RESOLUTION GENOME SCANNING
                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
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Pred. No. 1.6e+02;
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Pred. No. 1
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.6e+02;
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Best Local (
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CURRENT APPLICATION NUMBER: US/08/522,384
CURRENT FILING DATE: 1996-11-15
NUMBER OF SEQ ID NOS: 122
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 63
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                                             FILING DATE: October 26, 1995
ATTOREY AGENT INFORMATION:
NAME: WARDURG, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: LOPEZ-NIETO, CARLOS E APPLICANT: NIGAM, SANJAY KUMAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Unknown Organism
FEATURE:
INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                            CURRENT APPLICATION NUMBER FILING PARE: WORD BATA:
                                                                                                                                                             CLASSIFICATION: 514
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
     TELEFAX: (210)
               TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 633 West Fifth Street STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        l Similarity
7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   McSwiggen, James
Stinchcomb, Dan T.
Escobedo, Jaime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Application US/08584040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pavco, Pamela
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       METHOD AND REAGENT FOR THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           32.0%;
87.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TREATMENT OF DISEASES OR CONDITIONS RELATED TO LEVELS OF VASCULAR ENDOTHELIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GROWTH FACTOR
                                                                                                                                                                 60/005,974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8502
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Pred. No. 54;
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US-08-584-040-7661
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR FILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
SEQ ID NO 3450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 32.0%;
Best Local Similarity 87.5%;
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                      Sequence 34,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: RNA
ORGANISM: Mus sp
-09-371-772B-3450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 3450, Application US/09371772B Patent No. 6566127 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                           -07-949-541A-34/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00,876-J (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT FILING DATE: 1999-08-10
CURRENT FILING DATE: 1999-08-10
                                                                                                                                                                                                                                                                                                                     GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Ribozyme Pharmaceuticals, Inc
                                                                                                                                                                                                                                                                                                                                        tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ENGTH: 17
                                                                                                                 ent No. 5552270
NUMBER OF SEQUENCES:
                                                                                                    CORRESPONDENCE ADDRESS:
                                                                                                                                                        TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: sing
TOPOLOGY: linear
                               CITY: New York
STATE: New York
                                                                   STREET:
                  COUNTRY:
                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                     INFORMATION:
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10023
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Stinchcomb, Dan
Escobedo, Jaime
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                                                                                                                                                                                                                                                                                                                                                      Application US/07949541A
                                                                   26 West 61st Street
                                                                                                                                                                      Mirzabekov, Andrei D.
VENTION: Method for Determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                             Khrapko, Konstantin R. Khorlin, Alexandr A.
                                                                                                                                                                                                      Florentiev, Vladimir L.
                                                                                                                                                                                                                                              Ivanov, Igor B.
Ershov, Gennady M.
                                                                                                                                                                                                                               Lysov, Jury P
                                                                                      Ladas & Parry
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87.5%;
                                                                                                                                                        Sequence and a Device
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Pred. No. 66;
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Pred. No. (
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Query Match
Best Local Similarity
Thehes 6; Conserv:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                           APPLICANT: Florentic
APPLICANT: Mirzabeko
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: chemically synthesized MOLECULE TYPE: desoxyribooligonucleotide. FEATURE: oligonucleotide was synthesized by phosphoroamidite FEATURE: method.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 33,778
REFERENCE/DOCKET NUMBER: U-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 708-1800
TELEFAX: (212) 246-8959
                                                                     ZIP: 10023
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 8 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 18-Mar-1992
APPLICATION NUMBER: Russian Federation 4919321
FILING DATE: 18-Mar-1991
ATTORNEY/AGENT INFORMATION:
NAME: Janet I. Cord
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
COMPUTER: IBM PC/XT/AT OR OPERATING SYSTEM: DOS
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
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                                                                                                                                                                  STREET: 26 West 61st Street CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Diskette, 5.25 inch, 360 COMPUTER: IBM PC/XT/AT or compatibles
                                                                                                                              COUNTRY:
                                                                                                                                                                                                    ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                          5552270
                                                                                                                                               New York
                                                                                                                                                                                                                                                                           Florentiev, Viau....
Mirzabekov, Andrei D.
Mirzabekov Method for De
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/07949541A
                                                                                                                                USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                   Lysov,
                                                                                                                                                                                                                                                                                                                                                                                     Ivanov,
                                                      E: Diskette, 5.25 inch, 360
IBM PC/XT/AT or compatibles
                                                                                                                                                                                                      Ladas & Parry
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                                                                                                                                                                                                                                                                                                                                                                  Gennady M.
                                                                                                                                                                                                                                                                                                                                                                                       Igor B.
                                                                                                                                                                                                                                                                           Method for Determining a DNA Nucleotide
Sequence and a Device for Carrying Out Same
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30.0%; Score 6; DB 1; Le 100.0%; Pred. No. 1.6e+02;
                                                                                                                                                                                                                                           47
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                                                                                                                                                                                                                                                                                                                            Vladimir L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       34:
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                                                                       360 Kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kb storage
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US-08-593-345B-13/c
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PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: PCT/RU92/00052
FILING DATE: 18-Mar-1992
APPLICATION NUMBER: Russian Federation 4919321
FILING DATE: 18-Mar-1991
ATTORNEY/AGENT INFORMATION:
NAME: Janet I. Cord
NAME: Janet I. Cord
NAME: 33,778
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-07-949-541A-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                           COMPUTER: Macintosh
OPERATING SYSTEM: Macintosh 7.1
SOFTWARE: Wordporfect.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/593,345B
FILING DATE: 29-JAN-96
PRIOR APPLICATION DATA: No. 5851772e
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: chemically synthesized
MOLECULE TYPE: desoxyribooligonucleotide.
FEATURE: oligonucleotide was synthesized by phosphoroamidite
FEATURE: method.
OTHER INFORMATION: letter M means the tethering to the
OTHER INFORMATION: support via oxidized 3-methyluridine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 33,778
REFERENCE/DOCKET NUMBER: U-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 708-1800
NAME: Cherekov, Michael J.
REGISTRATION NUMBER: 33,664
REFERENCE/DOCKET NUMBER: ANL-IN-95-029+30
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 621-1330
                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: 3.50 inch, 1.4 MB storage
                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Dubiley, Svetlana A

TITLE OF INVENTION: A Microchip Method for the Enrichment

Specific DNA Sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELLEFAX: 122
TELEFAX: 233288
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                                                                                                                                                                                                                                                                                                                                       CITY: Chicago
STATE: Illinois
                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                              STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                   United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Lysov, Yuriy P
Shick, Valentine V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                CHERSKOV & FLAYNIK
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; Pred. No. 1.6e+02;
0; Mismatches 0;
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Matches 6; Conserv
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                                                                                                               TELEFAX: 713/651-5246
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tent No.
            MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "oligonucleotide" HYPOTHETICAL: YES
                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/859,954
ANTI-SENSE:
                                                                                                                                                                                   TELEPHONE: 713/651-5325
                                                                                                                                                                                                                                       FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Paul, Thomas D.
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Hardin, Paul E.
TITLE OF INVENTION: Gene Sequencing and Method Thereof
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U.S.A. ZIP: 77010-3095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IDENTIFICATION METHOD: Similarity with known sequences OTHER INFORMATION: Complementarity with primer of OTHER INFORMATION: exons to a-thalassemia gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: No. 5851772 Applicable
TOPOLOGY: linear
                                                                                STRANDEDNESS: single
                                                                                                  TYPE: nucleic acid
                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE:
                                                                FOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OCATION:
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1301 McKinney, Suite 5100
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Hardin, Paul E.
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100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            566
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o. 1.6e+02;
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US-08-859-954-53

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Query Match
Best Local Similarity
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                                                                               RESULT 79
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US-08-859-954-81
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US-08-859-954-81/c
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                    Sequence 104, Application US/08859954 Patent No. 6083695 GENERAL INFORMATION:
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Sequence 81, App...
Sequence 81, App...
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INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                                                 DESCRIPTION YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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APPLICANT: Homayouni, Ramin
APPLICANT: Hardin, Paul E.
                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Paul, Thomas D.
REGISTARTION UNMBER: 32,714
REFERENCE/DOCKET NUMBER: D-5900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                              MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "oligonucleotide"
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MEDIUM TYPE: Floppy
APPLICANT: Hardin, Susan H.
                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: single
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CITY: Houston
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/859,954
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                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
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77010-3095
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                                                                                                                                                                                                                                                                                                               Patent No. 6083695
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APPLICANT: Hardin, Susan H.
APPLICANT: Homayouni, Ramin
APPLICANT: Hardin, Paul E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
CLASSIFICATION:
PRIOR APPLICATION NUMBER: 08/63
APPLICATION NUMBER: 08/63
AFILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: PAUL, Thomas D.
REGISTRATION NUMBER: 32,7
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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REFERENCE/DOCKET NUMBER: D-5
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713/651-5325
TELEPAX: 713/651-5246
IFORMATION: TOT 707
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/859,954
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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TITLE OF INVENTION: Design
TITLE OF INVENTION: Gene St
TUTHER OF SEQUENCES: 566
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                                                                                           CITY: Houston
STATE: Texas
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                                                        COUNTRY: U.S.A. ZIP: 77010-3095
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                                                                                                                                                    ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                    329, Application US/08859954
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1301 McKinney, Suite 5100
                                                                                                                             E: Fulbright & Jaworski L.L.P
1301 McKinney, Suite 5100
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Gene Sequencing and Method Thereof
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100.0%;
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Pred. No.
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APPLICATE:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Paul, Thomas D.
REGISTRATION NUMBER: 32,714
REGISTRATION NUMBER: D-5900
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 107681-5325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 3 Patent No.
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Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                        COUNTRY: U.S.A.
ZIP: 77010-3095
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
OPTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/859,954
                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Fulbright & Jaworski L.L.P.
STREET: 1301 McKinney, Suite 5100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: other nucleic acid DESCRIPTION: /desc = "oligonucleotide" HYPOTHETICAL: YES
                                                                                                                                         FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Homayouni, Ramin
APPLICANT: Hardin, Paul E.
TITLE OF INVENTION: Design and Optimized Primer Library for
TITLE OF INVENTION: Gene Sequencing and Method Thereof
NUMBER OF SEQUENCES: 566
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NAME: Paul, Thomas D.
REGISTRATION NUMBER: 32,
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APPLICATION NUMBER:
FILING DATE:
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CITY: Houston
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ER: D-9
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                                                   ; ANTI-SENSE: US-08-859-954-333
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Best Local Similarity 10v.
                                                                                                                                                TELEPHONE: 713/651-5325
TELEPAX: 713/651-5246
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 8 base pairs
TYPE: nucleic acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Patent No. 6005000:
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 333,
                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/632,78
FILING DATE:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Paul, Thomas D.
REGISTRATION NUMBER: 32,714
REFERENCE/DOCKET NUMBER: D-59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 8 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: U.S.A.
ZIP: 77010-3095
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
COMPUTER: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "oligonucleotide"
HYPOTHETICAL: YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Homayouni, Ramin
APPLICANT: Hardin, Paul E.
TITLE OF INVENTION: Design and Optimized Primer Library for
TITLE OF INVENTION: Gene Sequencing and Method Thereof
NUMBER OF SEQUENCES: 566
                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION: TELEPHONE: 713/651-5325
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "oligonucleotide"
                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
CLASSIFICATION:
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/859,954
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Houston
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1301 McKinney, Suite 5100
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                                                                   YES
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                                                                                                                                       linear
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30.0%;
100.0%;
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Score 6;
Pred. No.
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o. 1.6e+02;
DB 1; Length 8; . 1.6e+02;
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US-08-859-954-360
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                                                     Sequence 396, Application US/08859954
Patent No. 6083695
GENERAL INFORMATION:
                                                                                                                                                                                                                                Query Match
Best Local Similarity
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                        APPLICANT:
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FILING DATE:
ATTORNEY/AGENT INFORMATION:
ATTORNEY Paul, Thomas D.
32,714
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: other nucleic acid

DESCRIPTION: /desc = "olgionucleotide"

HYPOTHETICAL: YES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCE ADDRESS:
CORRESPONDENCE ADDRESS:
Fulbright & Jaworski L.L.P.
ADDRESSEE: Fulbright & Jaworski L.L.P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Homayouni, Ramin
APPLICANT: Hardin, Paul E.
TITLE OF INVENTION: Design and Optimized Primer Library for
TITLE OF INVENTION: Gene Sequencing and Method Thereof
          APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: D-
TELECOMMUNICATION INFORMATION:
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COUNTRY: U.S.A.
ZIP: 77010-3095
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                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                             8 base pairs
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Homayouni, Ramin
Hardin, Paul E.
                                      Hardin, Susan H.
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Search completed: November 10, 2003, 08:03:59 Job time : 1 secs
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Best Local Similarity
Thehas 6; Conserve
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; ANTI-SENSE: 1
US-08-859-954-396
                                                                                                                                                                                                                                                                                                                                   TELEFAX: 713/651-5246
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 base pairs
                                                                                                                                                                                                                                                 MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "oligonucleotide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US
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TITLE OF INVENTION: Ge
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: 713/651-5325
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STALL COUNTRY: U.U.S. 77010-3095
                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Paul, Thomas D.
REGISTRATION NUMBER: 32
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                     TOPOLOGY:
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                                                                                                                                    Conservative
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Gene Sequencing and Method Thereof
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Maximum DB
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Maximum Match 100%
Listing first 86 summaries
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Perfect score:
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       Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GenCore version 5.1.6 (c) 1993 - 2003 Compugen Ltd.
                 US-09-774-809-31
US-09-774-809-42
US-09-888-326-43
US-09-888-326-43
US-09-78-479-313
US-09-78-479-313
US-10-117-95-311
US-10-007-010-56
US-10-007-010-56
US-10-30-627-1410
US-09-263-959-437
US-10-005-728-15
US-09-263-959-437
US-10-005-728-15
US-09-263-959-437
US-10-007-010-56
US-10-330-627-128
US-10-330-627-128
US-10-330-627-128
US-10-330-627-123
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                 Sequence 31, Appl Sequence 42, Appl Sequence 463, Appl Sequence 311, App Sequence 311, App Sequence 56, Appl Sequence 1410, App Sequence 1411, App Sequence 7, Appl Sequence 7, Appl Sequence 15, Appl Sequence 41, App Sequence 421, App Sequence 421, App Sequence 421, App Sequence 422, App Sequence 422, App Sequence 123, App Sequence 1231, App Sequence 1234, App Sequence 1234, App Sequence 1234, App Sequence 1234, App Sequence 1237, App Sequence 123
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quence 2144 quence 2144 quence 2144 quence 2144 quence 2167 quence 167 quence 167 quence 1211 quence 1211 quence 239 quence 239 quence 239 quence 239 quence 239 quence 239 quence 239 quence 239 quence 239 quence 1757 quence 1757 quence 1757	equence 123 equence 621 equence 127 equence 127 equence 127 equence 62, equence 62, equence 63, equence 153 equence 153 equence 153 equence 175	quence 2424 quence 2374 quence 2376 quence 2376 quence 2424 quence 2377 quence 2377

ALIGNMENTS

RESULT 1

US-09-774-809-31

US-09-774-809-31

Sequence 31, Application US/09774809

publication No. US20030004120A1

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: McKay, Robert A.
APPLICANT: Dean, Nicholas M.
APPLICANT: Monia, Brett
APPLICANT: Monia, Brett
APPLICANT: Morio, Pam
APPLICANT: Morio, William A.
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE COMPOSITIONS AND METHODS
TITLE OF INVENTION: FOR THE MODULATION OF JNK PROTEINS
FILE REFERENCE: ISPH-0412
CURRENT APPLICATION NUMBER: US/09/774,809

CURRENT FILING DATE: 2001-01-31

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT FILING DATE: 2001-01-31
PRIOR APPLICATION NUMBER: 09/396,902
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: 09/130,616
PRIOR FILING DATE: 1998-08-07
PRIOR APPLICATION NUMBER: 08/910,629
PRIOR FILING DATE: 1997-08-03
  Sequence 463, Application US/09888326
Publication No. US20030026801A1
GENERAL INFORMATION:
APPLICANT: Weiner, George
APPLICANT: Hartmann, Gunther
TITLE OF INVENTION: Methods for Enhancing Antibody-Induced
TITLE OF INVENTION: Cell Lysis and Treating Cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 165
SEQ ID NO 42
LENGTH: 20
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Publication No. US20030004120A1
GENERAL INFORMATION:
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PRIOR FILING DATE: 1997-08-03
NUMBER OF SEQ ID NOS: 165
SEQ ID NO 31
LENGTH: 20
TYPE: NO 30
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PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: 09/130,616
PRIOR FILING DATE: 1998-08-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/774,809
CURRENT FILING DATE: 2001-01-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Nero, Pam
APPLICANT: Gaarde, William A.
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE COMPOSITIONS AND METHODS
TITLE OF INVENTION: FOR THE MODULATION OF JNK PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: McKay, Robert A.
APPLICANT: Dean, Nicholas M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILE REFERENCE: ISPH-0412
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
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REFERENCE: C1049/7052 (AWS)
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Nero, Pam
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100.0%; Pred. No.
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           Sequence 301, Application US/10112653
Publication No. US20030050268A1
GENERAL INFORMATION:
APPLICANT: Krieg, Arthur M.
APPLICANT: Berg, Daniel J.
APPLICANT: Berg, Daniel J.
APPLICANT: OF INVENTION: IMMUNOSTIMULATORY NUCLEIC ACID FOR TITLE OF INVENTION: TREATMENT OF NON-ALLERGIC INFLAMMATORY DISEASES FILE REFERENCE: C01039/70060(AWS)
CURRENT APPLICATION NUMBER: US/10/112,653
CURRENT FILING DATE: 2002-03-29
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Matches
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 311
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 463
LENGTH: 20
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Best Local
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CURRENT APPLICATION NUMBER: US/09/776,479
CURRENT FILING DATE: 2001-02-02
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CURRENT FILING DATE: 2001-06-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/179,991 PRIOR FILING DATE: 2000-02-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Bratzler, Robert L.
APPLICANT: Petersen, Deanna M.
APPLICANT: Fouron, Yves
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PRIOR APPLICATION NUMBER: US 60/279,642
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LOCATION: (0)...(0)
OTHER INFORMATION: phosphorothioate backbone
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                   Local
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Pred. No. 0.
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Pred. No. 0.076;
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; OTHER INFORMATION: Synthetic Oligonucleotide
US-10-112-653-301
                                                      ; OTHER INFORMATION: Antisense Oligonucleotide US-10-007-010-56
                                                                                                                                                                                                                                                                                                                                                               US-10-007-010-56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: Synthetic Sequence US-10-017-995-311
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PRIOR FILING DATE: 2001-12-18
PRIOR FILING DATE: 2000-12-14
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FASESEQ for Windows Version 3.0
SEQ ID NO 311
LENGTH: 20
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Best Local S
Matches 20
                                                                                                                                                                                                                                    Sequence 56, Application US/10007010
Publication No. US20030125275A1
GENERAL INFORMATION:
APPLICANT: Alexander H. Borchers
APPLICANT: Kenneth W. Dobie
TITLE OF INVENTION: ANTISENSE MODULATION OF HCK EXPRESSION
FILE REFERENCE: RTS-0345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 311, Application US/10017995 Publication No. US20030055014A1 GENERAL INFORMATION:
                                                                                                         SEQ ID NO 56
LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
 Query Match
Best Local Similarity
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Best Local Similarity
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SOFTWARE: FRANKEQ for Windows Version 3.0
SEQ ID NO 301
LENGTH: 20
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CURRENT FILING DATE: 2001-12-04
NUMBER OF SEQ ID NOS: 87
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TITLE OF INVENTION: Inhibition of Angiogenesis by Nucleic Acids
FILE REFERENCE: C1037/7025 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/10/017,995
CURRENT FILING DATE: 2001-12-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 2001-03-29
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100.0%; Pred. No. 0.
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Score 17;
Pred. No.
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 DB 1;
0.59;
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                 Length 20;
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Best Local Similarity
Matches 10; Conserva
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                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1410
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 6
LENGTH: 17
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APPLICANT: Elledge, Stephen J.
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CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: US 09/448,480 PRIOR FILING DATE: 1999-11-24
                                                                                                                                                                                                                                                                                                             APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US/09/488,364
PRIOR FILING DATE: 2000-01-12
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CURRENT FILING DATE: 2001-12-12
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TITLE OF INVENTION: MAMMALIAN CHECKPOINT GENES AND PROTEINS
FILE REFERENCE: 120541-1013
                                                                                                                                                       TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
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                                                                                                                                                                    LENGTH:
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10 TCCGGGCCAG 1
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RESULT 10 US-10-330-627-1411/c · <aruence 1411, Application US/10330627

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APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1411
LENGTH: 10
Query Match
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                                                                                              US-09-263-959-437
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 12 base pairs
TYPE: nucleic acid
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ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: McMasters, David, D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 920010.426C2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC COMPOSITIONS AND METHODS WHICH UTI NUMBER OF SEQUENCES: 1279
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                          STRANDEDNESS: sir
TOPOLOGY: linear
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                    Conservative
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(206) 682-6031
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Koop, Ben F.
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                                    50.0%; Score 10; DB 1; Length 12; 100.0%; Pred. No. 11;
                    Mismatches
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CURRENT APPLICATION NUMBER: US/10/005,212
CURRENT FILING DATE: 2001-12-04
PRIOR APPLICATION NUMBER: USSN 08/986,997
PRIOR FILING DATE: 1997-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 7
LENGTH: 13
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APPLICANT: GILLIES, Stephen D
APPLICANT: LAN, Yan
APPLICANT: LO, Kin-Ming
TITLE OF INVENTION: Inmune Therapy and General Immune Stimulation
FILE REFERENCE: LEX-002C1
                                                                                                                                                                              SOFTWARE: PatentIn version 3.0
SEQ ID NO 1619
LENGTH: 10
                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: GENZYME CORPORATION
APPLICANT: ROBERTS, BRUCE
APPLICANT: SHANKARA, SRINIVAS
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Publication No.
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CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: PCT/US99/13800
PRIOR FILING DATE: 1999-06-18
NUMBER OF SEQ ID NOS: 2137
                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: PREPARATION AND USE OF FILE REFERENCE: GA0201C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                   TYPE: DNA
10 AGGCCAAAG 2
                             10 AGGCCAAAG 18
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No. US20020193570A1
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Pred. No. 16;
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RESULT 14 US-10-055-728-15/c

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; OTHER INFORMATION: TAG sequence Hs74649
US-10-055-728-15
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SEQ ID NO 15
LENGTH: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: US ZIP: 98104-7092
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
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PRIOR FILING DATE: 2001-09-28
PRIOR APPLICATION NUMBER: EP 0120373.2
PRIOR FILING DATE: 2001-09-28
PRIOR APPLICATION NUMBER: EP 01200228.3
PRIOR FILING DATE: 2001-01-23
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TITLE OF INVENTION: MEAKS AND METHODS FOR TREATMENT EVALUATION
FILE REFERENCE: 5244US (REN/P55190US00)
CURRENT APPLICATION NUMBER: US/10/055,728
CURRENT FILING DATE: 2002-04-19
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NUMBER OF SEQ ID NOS: 156
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                                    TELEFAX: (206) 682-6031 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                        SUFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/263,959
FILING DATE: 05-MAR-1999
CLASCIFY----
                                                                                                                                                   NAME: MCMASters, David D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 920010.426C2
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
ENGTH:
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12 base pairs
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Koop, Ben F.
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100.0%;
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FEATURE:
OTHER INFORMATION: Synthetic oligonucleotide
US-10-329-465-204
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                                                                                                                                                                                                                     SOFTWARE: PatentIn version 3.1 SEQ ID NO 204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/007,010
CURRENT FILING DATE: 2001-12-04
NUMBER OF SEQ ID NOS: 87
SEQ ID NO 56
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Best Local
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/10/329,465
CURRENT FILING DATE: 2002-12-23
                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                  APPLICANT: Wang et al.
TITLE OF INVENTION: GENES ABNORMALLY EXPRESSED IN MYELOID LEUKEMIA CELLS WITH AN MLL-1
TITLE OF INVENTION: FUSION
FILE REFERENCE: 27373/37928A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: ANTISENSE MODULATION OF HCK EXPRESSION FILE REFERENCE: RTS-0345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Alexander H. Borchers APPLICANT: Kenneth W. Dobie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                         TYPE: DNA
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                                                                                                                                                                                                      LENGTH:
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83.3%;
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Pred. No. 17
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Pred. No. 46;
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Pred. No. 23;
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                                                                                            DB 1;
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RESULT 18

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TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 420
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US-10-330-627-422
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US-10-330-627-421
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PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 421
LENGTH: 10
                                                                                                                                      Sequence 422, Application US/10330627 Publication No. US20030175771A1 GENERAL INFORMATION:
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APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE<sub>6</sub>: 2002-12-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OP INVENTION: Human Transcriptomes
FILE REFERENCE: 000107.00319
CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
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Local Similarity 90.0%;
les 9; Conservative
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les 9; Conserv
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Pred. No. 1
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; ORGANISM: Homo sapiens US-10-330-627-1228
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Best Local Similarity
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US-10-330-627-644
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APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
                                                                                                                                                                                                                                                                                    Sequence 1228, Application US/10330627
Publication No. US20030175771A1
GENERAL INFORMATION:
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 644
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 422
                                                                       SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 1228
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Publication No. US20030175771A1
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Best Local Similarity
                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/330,627 CURRENT FILING DATE: 2002-12-30
                                                                                                                                                                                              APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
                                                                                                                            PRIOR APPLICATION NUMBER: US 09/448,480 PRIOR FILING DATE: 1999-11-24
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                                                                                                         NUMBER OF SEQ ID NOS: 1564
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                                   TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Homo sapiens
                                                  LENGTH: 10
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Pred. No. 17
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Pred. No. :
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Query Match

42.0%; Score 8.4;

DB 1;

Length 10;

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Sequence 1230, Application US/10330627
Publication No. US20030175771A1
GENERAL INFORMATION:
APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
RESULT 25
US-10-330-627-1231/c .
; Sequence 1231, Application US/10330627
                                                                                                                                                                                                                                                ; ORGANISM: Homo sapiens US-10-330-627-1230
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1230
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PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1229
LENGTH: 10
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CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
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APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
                                                                                                                                                                                                                                                                                 LENGTH: 10
TYPE: DNA
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                                                                                                                                                                     / Match 42.0%;
Local Similarity 90.0%;
les 9; Conservative
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les 9; Conser
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90.0%;
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Sequence 1233, Application US/10330627
Publication No. US20030175771A1
GENERAL INFORMATION:
APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US/0/448,480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1232
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; ORGANISM: Homo sapiens
US-10-330-627-1232
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US-10-330-627-1232/c
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-330-627-1231
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CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1231
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GENERAL INFORMATION:
APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
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Best Local Similarity 90.0%;
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APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
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Pred. No. 17;
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Pred. No. 1
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FILE REFERENCE: 001107.00319
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 1234
LENGTH: 10
TYPE: DNA
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US-10-330-627-1233
                                                                                                                                                US-10-033-145-405
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1233
                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 2137
SOFTWARE: PatentIn version 3.0
SEQ ID NO 405
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Best Local Similarity Matches 9; Conserva
                                                                        Query Match
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PRIOR FILING DATE: 1999-06-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: GENZYME CORPORATION
APPLICANT: ROBERTS, BRUCE
APPLICANT: SHANKARA, SRINIVAS
TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/10/033,145
CURRENT FILING DATE: 2001-11-05
                                                                                                                                                                             ORGANISM: Homo sapiens
                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                                                                                                              ENGTH: 10
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9; Conserv
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9; Conservative
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Vogelstein, Bert
   Conservative,
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90.0%;
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; Score 8.4; DB ; Pred. No. 17; 0; Mismatches
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Pred. No.
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PRIOR APPLICATION NUMBER: PCT/US99/13800
PRIOR FILING DATE: 1999-06-18
NUMBER OF SEQ ID NOS: 2137
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1326
LENGTH: 10
TYPE: DNA
ORGANISM: Homo sapiens
US-10-033-145-1326
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                                                                                       Query Match
Best Local
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APPLICANT: Cornelissen, Marion
TITLE OF INVENTION: MEANS AND METHODS FOR TREATMENT EVALUATION
FILE REFERENCE: 5244US (REN/P55190US00)
CURRENT APPLICATION NUMBER: US/10/055,728
CURRENT FILING DATE: 2002-04-19
PRIOR APPLICATION NUMBER: 60/325,722
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CURRENT FILING DATE: 2001-11-05
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TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
FILE REFERENCE: GA0201C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: GENZYME CORPORATION APPLICANT: ROBERTS, BRUCE
                                                                                                                                                                                                                                                                                                                                                                 PRIOR FILING DATE: 2001-09-28
PRIOR APPLICATION NUMBER: EP 0120373.2
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                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: EP 01200228.3
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                                                                                                                                                                                               TYPE: DNA ORGANISM: Artificial Sequence
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                                                                        Similarity 9; Conserv
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Pred. No. 23;
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; OTHER INFORMATION: DNA
US-09-989-789-2376
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Sequence 2424, Application US/09989789
Patent No. US20020063379A1
GENERAL INFORMATION:
APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/09/989,789
CURRENT FILING DATE: 2002-03-25
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NUMBER OF SEQ ID NOS: 4085
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 2376
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APPLICANT: LIU, Qiang
APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/09/989,789
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Patent No. US20020063379A1
GENERAL INFORMATION:
APPLICANT: LIU, Qiang
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Best Local
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA ORGANISM: Artificial Sequence FEATURE:
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Pred. No.
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US-99-990-186-2376/c

Sequence 2376, Application US/09990186

Publication No. US20030068675A1

GENERAL INFORMATION:

APPLICANT: LIU, Qiang

TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS

FILE REFERENCE: 8325-0011.21 / S11-US3

CURRENT APPLICATION NUMBER: US/09/990,186

CURRENT APPLICATION NUMBER: US/09/990,186

CURRENT FILING DATE: 2001-11-20

NUMBER OF SEQ ID NOS: 4085

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 2376
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Matches
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TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.21 / S11-US3
CURRENT APPLICATION NUMBER: US/09/990,186
CURRENT FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2374
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Best Local :
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FEATURE:
OTHER INFORMATION:
OTHER INFORMATION:
                                                              TYPE: DNA
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                                            ORGANISM: Artificial Sequence
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Description DNA
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Pred. No.
                of Artificial Sequence:
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RESULT 38
US-09-989-994-2374/c
US-09-989-994-2374, Application US/09989994
Publication No. US20030104526A1
GENERAL INFORMATION:
GENERAL INFORMATION:
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RESULT 39
US-09-989-994-2376/c
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Publication No. US20030068675A1
GENERAL INFORMATION:
APPLICANT: LIU, Qiang
                                                                                                                                                                                                                                                                            Query Match
Best Local
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Best Local :
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CURRENT FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
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CURRENT APPLICATION NUMBER: US/09/990,186
CURRENT FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
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8; Conservation
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Sequence 1228, Application US/10330627
Publication No. US20030175771A1
GENERAL INFORMATION:
APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
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Best Local S
Matches 8
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Publication No. US20030104526A1
GENERAL INFORMATION:
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CURRENT FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 2376
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Best Local
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TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
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TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
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ORGANISM: Artificial Sequence
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PRIOR APPLICATION NUMBER: US 09/448,480

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Sequence 1230, Application US/10330627
Publication No. US20030175771A1
GENERAL INFORMATION:
APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR FILING DATE: 1999-11-24
PRIOR FILING DATE: 1999-11-24
SOPTWARE: FASHESEQ for Windows Version 4.0
SEQ ID NO 1230
LENGTH: 10
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Publication No. US20030175771A1

GENERAL INFORMATION:
APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W.
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1229
LENGTH: 10
                                                                                                                       ; ORGANISM: Homo sapiens US-10-330-627-1230
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US-10-330-627-1230
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NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FBATSEQ for Windows Version 4.0
SEQ ID NO 1228
Query Match
Best Local Similarity
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Local Similarity 100.0%; Find the second second
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les 8; Conserv
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0; Mismatches
Score 8; DB 1;
Pred. No. 22;
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Pred. No.
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5. 22;
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FILE REFERENCE: 001107.00319
CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1232
LENGTH: 10
TYPE: DNA
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APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT FAPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1231
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Best Local Similarity
Watches 8; Conserva
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US-10-330-627-1231
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                                                               Best Loc
Matches
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APPLICANT: Velculescu, Victor E.
                                                                             Query Match
Best Local
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APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
                                                                                                                                              ORGANISM: Homo sapiens
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100.0%; Pred. No.
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100.0%; FI
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100.0%; Pred. No.
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RESULT 46
US-10-330-627-1233
; Sequence 1233, Application US/10330627
; Publication No. US20030175771A1

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APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1234
LENGTH: 10
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1233
LENGTH: 10
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Best Local (
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                                                        APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
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CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24'
                      PRIOR APPLICATION NUMBER: US 09/448,480 PRIOR FILING DATE: 1999-11-24
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APPLICANT: Kinzler, Kenneth
APPLICANT: Vogelstein, Bert
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SEQ ID NOS: 1564
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k; Pred. No. 22;
0; Mismatches
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0; Mismatches
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Query Match
Best Local Similarity
Thes 8; Conserve
Query Match
Best Local Similarity
Thes 8; Conserve
                                                                                     ; ORGANISM: Homo sapiens 
US-10-330-627-877
                                                                                                                                                                                                  APPLICANT: Veloulescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Kinzler, Kenneth W
APPLICANT: Kinzler, Kenneth W
APPLICANT: Kogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
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US-10-330-627-877/c
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Best Local Similarity
Thes 8; Conserv
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                                                                                                                                                  NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 877
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Publication No. US20030175771A1
GENERAL INFORMATION:
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 876
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Publication No. US20030175771A1
GENERAL INFORMATION:
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SEQ ID NO 621
LENGTH: 10
TYPE: DNA
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APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: US 09/448,480 PRIOR FILING DATE: 1999-11-24
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CURRENT FILING DATE: 2002-12-30
                                                                                                                   TYPE: DNA
                                                                                                                                    LENGTH: 10
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100.0%; Pred. No. 22
tive 0; Mismatches
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100.0%; Pred. No.
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APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1215
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US-10-330-627-1214/c
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                                                                                                                                                                                                                                                                                             ; ORGANISM: Homo sapiens
US-10-330-627-1215
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US-10-330-627-1214
                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.0%; Sc
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PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 1214
LENGTH: 10
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Publication No. US20030175771A1
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APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 10
TYPE: DNA
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Query Match
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                                                             ; ORGANISM: Homo sapiens US-10-033-145-62
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                                                                                                                                                                                                                                                                                                                                                                                                                                   US-10-033-145-62/c
                                                                                                               SOFTWARE: PatentIn version 3.0 SEQ ID NO 62 LENGTH: 10
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 62, Application US/10033145
Publication No. US20020151515A1
  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/033,145
CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: PCT/US99/13800
PRIOR FILING DATE: 1999-06-18
NUMBER OF SEQ ID NOS: 2137
                                                                                                                                                                                                                                                                            FILE REFERENCE: GA02010
                                                                                                                                                                                                                                                                                            APPLICANT: SHANKARA, SRINIVAS
TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
                                                                                                                                                                                                                                                                                                                                   APPLICANT: GENZYME CORPORATION APPLICANT: ROBERTS, BRUCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 10
                                                                                                 TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/07/882,885
FILING DATE: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 197/173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: METHOD AND REAGENT FOR INHIBITING P-GLYCOPROTEIN mdr-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
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CLASSIFICATION: <Unknown>
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STATE: California
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Score 8; E
Pred. No.
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Pred. No.
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GENERAL INFORMATION:
APPLICANT: GENZYME CORPORATION
APPLICANT: ROBERTS, BRUCE
APPLICANT: SHANKARA, SRINIVAS
ITILE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
FILE REFERENCE: GAO201C
CURRENT APPLICATION NUMBER: US/10/033,145
CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: PCT/US99/13800
PRIOR FILING DATE: 1999-06-18
UNMBER OF SEQ ID NOS: 2137
SOFTWARE: Patentin version 3.0
SEQ ID NO 1531
LENGTH: 10
                                RESULT 57
US-10-055-728-15
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Best Local Similarity
Whiches 8; Conserve
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US-10-033-145-1750
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US-10-033-145-1531
Sequence 15, Application US/10055728
Publication No. US20030170720A1
                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 2137
SOFTWARE: PATENTIN VETSION 3.0
SEQ ID NO 1750
LENGTH: 10
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Publication No. US20020151515A1
GENERAL INFORMATION:
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Sequence 1531, Application US/10033145
Publication No. US20020151515A1
                                                                                                                                                                              Query Match
Best Local 9
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CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: PCT/US99/13800
PRIOR FILING DATE: 1999-06-18
                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: GENZYME CORPORATION
APPLICANT: ROBERTS, BRUCE
APPLICANT: SHANKARA, SRINIVAS
TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
FILE REFERENCE: GA0201C
                                                                                                                                                                                                                                                           TYPE: DNA
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Best Local S
Matches 9
                                                            Sequence 31, Applica Publication No. US20 GENERAL INFORMATION:
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Best Local
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             APPLICANT: McKay, Robert A.
APPLICANT: Dean, Nicholas M.
APPLICANT: Monia, Brett
   APPLICANT:
                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                     LENGTH:
   Nero, Pam
                                                                                            Application US/09774809
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; FEATURE:
; OTHER INFORMATION: TAG sequence Hs74649
US-10-055-728-15
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CURRENT APPLICATION NUMBER: US/10/055,728
CURRENT FILING DATE: 2002-04-19
PRIOR APPLICATION NUMBER: 60/325,722
PRIOR FILING DATE: 2001-09-28
PRIOR APPLICATION NUMBER: EP 0120373.2
PRIOR FILING DATE: 2001-09-28
PRIOR FILING DATE: 2001-01-28
PRIOR FILING DATE: 2001-01-23
PRIOR FILING DATE: 2001-01-3
PRIOR FILING DATE: 2001-01-3
                                                                                                                                                                                                                                                                                                                                                                                               Sequence 6, Application US/10020038
Publication No. US20020156247A1
GENERAL INFORMATION:
APPLICANT: Elledge, Stephen J.
APPLICANT: Sanchez, Yolanda
TITLE OF INVENTION: MAMMALIAN CHECKPOINT GENES AND PROTEINS
FILE REFERENCE: 120541-1013
CURRENT APPLICATION NUMBER: US/10/020,038
CURRENT FILING DATE: 2001-12-12
                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn version 3.1
SEQ ID NO 15
LENGTH: 11
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                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US/09/488,364
PRIOR FILING DATE: 2000-01-12
NUMBER OF SEQ ID NOS: 15
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APPLICANT: Cornelissen, Marion
TITLE OF INVENTION: MEANS AND METHODS FOR TREATMENT EVALUATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence:
                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
11 GGCCTGTCCAA 1
                                         6 GGCCAGGCCAA 16
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                                                                                                      Similarity
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Pred. No. 53;
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Pred. No. :
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US20030004120A1

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FILE REFERENCE: ISPH-0412
CURRENT APPLICATION NUMBER: US/09/774,809
CURRENT FILING DATE: 2001-01-31
PRIOR APPLICATION NUMBER: 09/396,902
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: 09/130,616
PRIOR FILING DATE: 1998-08-07
PRIOR FILING DATE: 1998-08-07
PRIOR APPLICATION NUMBER: 08/910,629
PRIOR APPLICATION NUMBER: 08/910,629
PRIOR APPLICATION NUMBER: 08/910,629
PRIOR FILING DATE: 1997-08-03
NUMBER OF SEQ ID NOS: 165
SEQ ID NO 42
LENGTH: 20
RESULT 61
US-09-888-326-463/c
Sequence 463, Application US/09888326
Publication No. US20030026801A1
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CURRENT FILING DATE: 2001-01-31
PRIOR APPLICATION NUMBER: 09/396,902
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: 09/130,616
PRIOR APPLICATION NUMBER: 09/910,629
PRIOR APPLICATION NUMBER: 08/910,629
PRIOR APPLICATION NUMBER: 08/910,629
PRIOR FILING DATE: 1997-08-03
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SEQ ID NO 31
LENGTH: 20
TYPE: DNA
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Best Local Similarity
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Best Local Similarity
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Publication No. US20030004120A1
                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 20
TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Gaarde, william A.
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE COMPOSITIONS AND METHODS
TITLE OF INVENTION: FOR THE MODULATION OF JNK PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Gaarde, William A.
FITTLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE COMPOSITIONS AND METHODS
FITTLE OF INVENTION: FOR THE MODULATION OF JNK PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Monia, Brett
APPLICANT: Nero, Pam
                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence FEATURE:
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APPLICANT: Bratzler, Robert L.
APPLICANT: Petersen, Deanna M.
APPLICANT: Pouron, Yves
TITLE OF INVENTION: Immunostimulatory Nucleic Acids for the
TITLE OF INVENTION: Treatment of Asthma and Allergy
FILE REFERENCE: C1037/7013 (HCL/MAT)
CURRENT APPLICATION NUMBER: US/09/776,479
CURRENT FILING DATE: 2001-02-02
PRIOR APPLICATION NUMBER: US 60/179,991
PRIOR APPLICATION NUMBER: US 60/179,991
PRIOR APPLICATION NUMBER: US 60/179,991
PRIOR FILING DATE: 2000-02-03
NUMBER OF SEO ID NOS: 1093
SOFTWARE: FastSEQ for Windows Version 3.0
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 463
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                                                                              Matches
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Best Local Similarity
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Best Local
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APPLICANT: Hartmann, Gunther
TITLE OF INVENTION: Methods for Enhancing Antibody-Induced
TITLE OF INVENTION: Cell Lysis and Treating Cancer
FILE REFERENCE: C1039/7052 (AWS)
CURRENT APPLICATION NUMBER: US/09/888,326
CURRENT FILING DATE: 2001-06-22
PRIOR APPLICATION NUMBER: US 60/213,346
PRIOR FILING DATE: 2000-06-22
PRIOR FILING DATE: 2000-06-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -09-776-479-311/c
                                                                                                                                                                                                            LENGTH: 20
TYPE: DNA
ORGANISM: Artificial Sequence
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TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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14 GGCCTGGCCCGGAC 1
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Pred. No. 46;
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Pred. No. 4
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; Sequence 301, Applic ; Publication No. US20 ; GENERAL INFORMATION: ; APPLICANT: Krieg, A

Application US/10112653
o. US20030050268A1

US-10-112-653-301/c

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US-09-989-789-2148
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US-10-017-995-311/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION. Inhibition of Angiogenesis by Nucleic Acids FILE REFERENCE: C1037/7025 (HCL/MAT)
CURRENT APPLICATION UNMEBR: US/10/017,995
CURRENT FILING DATE: 2001-12-18
PRIOR APPLICATION UNMBER: US 60/255,534
PRIOR FILING DATE: 2000-12-14
PRIOR FILING DATE: 2000-12-14
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 311
LENGTH: 20
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                                 SOFTWARE: Pat
SEQ ID NO 2148
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                                                                                           APPLICANT: LIU, Qiang
APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
FILE REFERENCE: 8325-0011.20 / S19-789
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Best Local :
                                                                      CURRENT APPLICATION NUMBER: US/0 CURRENT FILING DATE: 2002-03-25 NUMBER OF SEQ ID NOS: 4085
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CURRENT APPLICATION NUMBER: US/10/112,653
CURRENT FILING DATE: 2002-03-29
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TYPE: DNA
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OTHER INFORMATION: Description of Artificial Sequence: example target; OTHER INFORMATION: DNA
US-09-989-789-2149
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Publication No. US20030068675A1
GENERAL INFORMATION:
APPLICANT: LIU, Qiang
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LENGTH: 9
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CURRENT FILLING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
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CURRENT APPLICATION NUMBER: US/09/
CURRENT FILING DATE: 2002-03-25
NUMBER OF SEQ ID NOS: 4085
                                                                                                                                                                                                                                                                               TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS FILE REFERENCE: 8325-0011.21 / S11-US3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
                                                                                                           OTHER INFORMATION:
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                                                                                                                                           FEATURE:
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88.9%;
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                                                          Score 7 4;
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Pred. No. :
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                                                                                                                          of Artificial Sequence: example target
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US-09-989-994-2148
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TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/09/989,994
CURRENT FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2148
                                                      Sequence 2149, Application US/09989994
Publication No. US20030104526A1
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SEQ ID NO 2149
LENGTH: 9
                                         GENERAL INFORMATION
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Publication No. US20030104526A1
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CURRENT APPLICATION NUMBER: US/09/990,186
CURRENT FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
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Pred. No. 1
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                                                                                   CURRENT APPLICATION NUMBER: US/10/033,145
CURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: PCT/US99/13800
PRIOR FILING DATE: 1999-06-18
NUMBER OF SEQ ID NOS: 2137
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SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2149
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                                                      SOFTWARE: PatentIn version 3.0 SEQ ID NO 405
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Best Local
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                                                                                                                                                                           APPLICANT: GENZYME CORPORATION
APPLICANT: ROBERTS, BRUCE
APPLICANT: SHANKARA, SRINIVAS
TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
FILE REPERENCE: GA0201C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Nucleic Acid Treatment of Diseases TITLE OF INVENTION: Epidermal Growth Factor Receptors FILE REFERENCE: MBHB00-958-K (400/064) CURRENT APPLICATION NUMBER: US/10/277,494 CURRENT FILING DATE: 2002-10-21
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APPLICANT: McSwiggen, Jim
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ORGANISM: Artificial Sequence
FEATURE:
ORGANISM: Homo sapiens
                    TYPE: DNA
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o. US20030186909A1
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88.9%;
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Pred. No. 1e+02;
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Pred. No. le+02;
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APPLICANT: Velculescu, Victor E.
APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
TITLE OF INVENTION: Human Transcriptomes
FILE REFERENCE: 001107.00319
CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1215
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CURRENT APPLICATION NUMBER: US/10/330,627
CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 1214
LENGTH: 10
TYPE: DNA
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US-10-330-627-1214
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US-10-330-627-1215
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US-10-330-627-1214
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Publication No. US20030175771A1
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APPLICANT: Kinzler, Kenneth W
APPLICANT: Vogelstein, Bert
                                                                                                                                                     LENGTH: 10
TYPE: DNA
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                              GCCAGGCCA 15
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GCCTGGCCA 9
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Pred. No.
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US-09-989-789-2396/c
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CURRENT FILING DATE: 2002-03-25
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: PATENTIN VEY: 2.0
SEQ ID NO 2396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
FILE REFERENCE: 8325-0011.20 / S11-US2
               Sequence 2397, Application US/09989789 Patent No. US20020063379A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/837,751
CURRENT FILING DATE: 2001-04-18
PRIOR APPLICATION NUMBER: US 60/198,124
PRIOR FILING DATE: 2000-04-18
NUMBER OF SEQ ID NOS: 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Green, Allan
APPLICANT: Singh, Surin
APPLICANT: Liu, Qing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Liu, Qing
TITLE OF INVENTION: Method of Modifying the Content of Cottonseed Oil
                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 9
TYPE: DNA
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                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: example target OTHER INFORMATION: DNA
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Pred. No.
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o. le+02;
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APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS

옃

GNN NUCLEOTIDE

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; Sequence 2396, Application US/09990186
; Publication No. US20030068675A1
; GENERAL INFORMATION:
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                                                        APPLICANT: LIU, Qiang
APPLICANT: LIU, Qiang
APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REPERENCE: 8325-0011.21 / S11-US3
CURRENT APPLICATION NUMBER: US/09/990,186
CURRENT APPLICATION NUMBER: US/09/990,186
CURRENT FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2397
LENGTH: 9
                                                                                                                                                                                                                                                     Sequence 2397, Application US/09990186 Publication No. US20030068675A1 GENERAL INFORMATION:
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Matches
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CURRENT FILING DATE: 2002-03-25
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 2397
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TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.21 / S11-US3
CURRENT APPLICATION NUMBER: US/09/990,186
CURRENT FILLING DATE: 2001-11-20
CURRENT FILLING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
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ORGANISM: Artificial Sequence
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OTHER INFORMATION:
                                     ORGANISM: Artificial Sequence
OTHER INFORMATION: Description of Artificial Sequence: example target
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les 7; Conserv
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100.0%; Pr
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Pred. No.
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Pred. No.
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b. le+02;
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GENERAL INFORMATION:

APPLICANT: LIU, Qiang
TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF
TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS
FILE REFERENCE: 8325-0011.20 / S11-US2
CURRENT APPLICATION NUMBER: US/09/989,994
CURRENT FILING DATE: 2001-11-20
NUMBER OF SEQ ID NOS: 4085
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 2396
LENGTH: 9
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                                                        Query Match
Best Local S
Matches 7
                                                                                                                                                                                                                      TITLE OF INVENTION: POSITION DEPENDENT RECOGNITION OF GNN NUCLEOTIDE TITLE OF INVENTION: TRIPLETS BY ZINC FINGERS FILE REPERENCE: 8325-0011.20 / S11-US2 CURRENT APPLICATION NUMBER: US/09/989,994 CURRENT FILING DATE: 2001-11-20 NUMBER OF SEQ ID NOS: 4085 SOFTWARE: Patentin Ver. 2.0 SEQ ID NO 2397
                                                                                                                                                                                                                                                                                                                                                Sequence 2397, Application US/09989994
Publication No. US20030104526A1
GENERAL INFORMATION:
APPLICANT: LIU, Qiang
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Best Local
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OTHER INFORMATION: Description
OTHER INFORMATION: DNA
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                                                                                                                                                               ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                             TYPE: DNA
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Pred. No.
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Pred. No.
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                                                          Mismatches
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o. le+02;
                                                                       DB 1;
. 1e+02;
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Publication NG. ...

Publication:

GENERAL INFORMATION:

APPLICANT: GENZYME CORPORATION

APPLICANT: NO SERTS, BRUCE

APPLICANT: SHANKARA, SRINIVAS

TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES

TITLE OF INVENTION: PREPARATION AND USE OF SUPERIOR VACCINES
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Best Local Similarity
Thes 8; Conserve
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US-10-330-627-877
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US-10-330-627-876
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 877
LENGTH: 10
                                                                                                                                                                         Sequence 1750, Application US/10033145 Publication No. US20020151515A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Velculescu, Victor E. APPLICANT: Kinzler, Kenneth W APPLICANT: Vogelstein, Bert
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CURRENT FILING DATE: 2002-12-30
PRIOR APPLICATION NUMBER: US 09/448,480
PRIOR FILING DATE: 1999-11-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/330,627
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION: Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 1564
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Homo sapiens
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Local Similarity 80.0%;
ses 8; Conservation
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CURRENT APPLICATION NUMBER: US/10/033,145
COURRENT FILING DATE: 2001-11-05
PRIOR APPLICATION NUMBER: PCT/US99/13800
PRIOR FILING DATE: 1999-06-18
NUMBER OF SEQ ID NOS: 2137
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1750
LENGTH: 10
TYPE: DNA
ORGANISM: Homo sapiens
US-10-033-145-1750
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                                                                                                                                                 RESULT 86
US-10-027-632-176277
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                                                                                                              Sequence 176277, Application US/10027632 GENERAL INFORMATION:
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Best Local
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Publication No. US20030204075A9
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Best Local
               APPLICANT: Wang, David G.

TITLE OF INVENTION: Identification and Mapping of Single Nucleotide TITLE OF INVENTION: Polymorphisms in the Human Genome FILE REFERENCE: 108827.129

CURRENT APPLICATION NUMBER: US/10/027,632
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
CURRENT FILING DATE:
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PRIOR APPLICATION NUMBER: US 60/198,676
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CURRENT FILING DATE: 2002-04-30
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PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-04-20
PRIOR FILING DATE: 2000-04-20
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-02-24
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PRIOR FILING DATE: 1000-02-24
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PRIOR FILING DATE: 1000-02-29
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PRIOR FILING DATE: 1000-02-38
PRIOR APPLICATION NUMBER: US 60/146,002
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PRIOR APPLICATION NUMBER: US 60/146,002
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PRIOR FILING DATE: 1000-02-38
PRIOR APPLICATION NUMBER: US 60/146,002
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                 Different Gene Expression Responses in Cotton Root and Hypocotyl tissues during infection with Fusarium Wilt Disease Unpublished (2003)

Contact: Caitriona Dowd, Helen McFadden Commonwealth Scientific and Industrial Research Organisation Division of Plant Industry Black Mountain Laboratories, Cnr Clunies Ross Street & Barry Driv Black Mountain, Canberra, ACT, 2601, Australia Tel: 61 2 6246 5914, 6246 5377
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8 bp mRNA linear EST 04-JUN-
CRH2.4F08 Cotton Root and Hypocotyl Lambda ZIPLOX Library (CRH)
Gossypium hirsutum cDNA clone CRH2.4F08 5', mRNA sequence.
CD486474
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Gossypium hirsutum
Gossypium hirsutum
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosid;
eurosids II; Malvales; Malvaceae; Malvoideae; Gossypium.
1 (bases 1 to 8)
Dowd, C., Wilson, I. and McFadden, H.
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Email: Caitriona.Dowd@csiro.au, Helen.McFadden@csiro.au
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Gossypium hirsutum cDNA clone CRH2.4F08 5', mRNA sequence.
CD486474
CD486474.1 GI:31407439
EST.
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Commonwealth Scientific and Industrial Research Organisation
Division of Plant Industry
                                                                                                                                                                                                                                                                                                                                                                                                                           Different Gene Expression Responses in Cotton Root and Hypocotyl tissues during infection with Fusarium Wilt Disease Unpublished (2003)
                                                                                                                                                                                                                        Email: Caitriona.Dowd@csiro.au, Vector clipped sequences Bases
                                                                                                                                                                                                                                                                    Black Mountain Laboratoriés, Cnr Clunies Ross Street & Barry Drive,
Black Mountain, Canberra, ACT, 2601, Australia
Tel: 61 2 6246 4914, 6246 5377
Fax: 61 2 6246 5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Malvales; Malvaceae; Malvoideae; Gossypium.
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Location/Qualifiers
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/organism="Gossypium hirsutum"
/mol type="mRNA"
/cultivar="DeltaEMERALD"
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/clone="CRH2.4F08"
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/dev_stage="5 day old seedlings"
/lab_host="Y1090(ZL)"
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Library (CRH)"
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/clone="CRH2.4F08"
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/mol_type="mRNA"
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Pred. No.
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/dev_stage="5 day old seedlings"
/lab_host="Y1090(ZL)"
/clone_lib="Cotton Root and Hypocotyl Lambda ZIPLOX
Library (CRH)"
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/mRNA was prepared from root and hypocotyl tissues of the
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phage vector (Life Technologies). Constructed by Caitriona
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